



bio-~~t~~echne®

SIMPLE WESTERN

基本原理及实验操作

王娴婷 FAS

Tel: (+86) 185-1660-6926

E-mail: xianting.wang@bio-techne.com

PROTEIN SIMPLE -- 创新蛋白质分析技术专家



Jess/Abby/Wes
全自动 Western



FluorChem
多功能成像



Ella
微流控全自动 ELISA



Milo
单细胞 Western

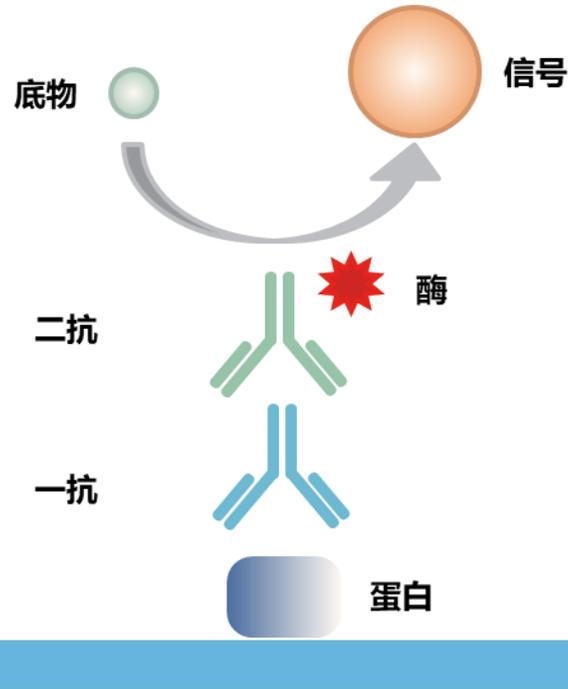


MFI
微流成像颗粒计数分类仪



Maurice
CE-SDS + icIEF 双功能 CE

免疫测定



典型的一抗结合目标蛋白的免疫测定。二抗结合一抗。二抗耦连酶，该酶能催化底物反应，产生能够检测的光信号。

- 任何使用抗体对混合蛋白中的靶分子或者目标蛋白进行检测的试验都叫**免疫测定**
 - 一些传统的免疫测定实验如：ELISAs、Western Blots.
- 一抗：检测目标蛋白
- 二抗：与一抗结合，辅助检测
 - 一般耦连酶或者显色基团
- 抗体复合物和目标蛋白可以多种方式检测
 - 化学发光（Chemiluminescence）
 - 荧光（Fluorescence）

当你在做 WESTERN BLOT 时



*图片来源于网络

当你搜索 WESTERN BLOT 时

豆 【转】五年western blot 实验经验总结

<https://www.douban.com/note/1364122221>

2011-2-24 · 4 Western Blot实验关键 Western

言, 扩
的选择

知 为什么同一批次提取

<https://www.biomedcentral.com/extra>

2013-8

知 血泪教训, Western blot史上最全避雷手册。 - * 乎

<https://zhuanlan.zhihu.com/p/100000000>

2020-9-3 · 提起Western blot

验, 简称“玄学”。话不多说, 顺利利。倾尽全力...

简 你的Western Blot

<https://www.jianshu.com/p/100000000>

2020-8-24 · 做Western Blot

昨天的差异这

知 [实验经验]组织样本的western blot, 这十点需要注意 - * 乎

<https://zhuanlan.zhihu.com/p/100000000>

知 如何解决western blot半

<https://wenda.guidechem.com/question/100000000>

2019-3-12 · 如何解决western blot半

显色的会影响条带的趋势, 已把ECL均匀的滴在整张膜上, 看见荧光后, 曝光, 得到一种趋势结果, 然后, 这张膜荧光渐渐黯淡, 为了增加强度, 重新再滴加ECL, 也覆盖的很均匀 ...

知 【求助】Western blot内参actin结果重复不出来 - 经验共享 ...

<https://bbs.antpedia.com/thread-228440-1-1.html>

2013-6-1 · 标题: 【求助】Western blot内参actin结果重复不出来 暗香涌[使用道具] 三级 UID 108870 精华

离线 1 发表于

知 Western Blot详解 - 常见的问题指南 (一) - 专区 - 生 * 谷

<https://news.bioon.com/article/6005931.html>

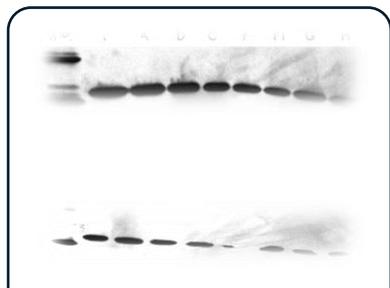
W. Western Blot 中抗体的重复应用问题 解答: 抗体工作溶液一般不主张储存反复使用, 但是如抗体比较珍贵, 可有使用? 2017 转载后应在? 2.5.1.1

知 Western Blot详解 - 常见的问题指南 (二) - 专区 - 生 * 谷

<https://www.bioon.com/article/6005934.html>

就重复性而言: 尼龙膜可反复用于分子杂交, 杂交后, 探针分子可经碱变性被洗脱下来; 硝酸纤维素膜不能重复使用; PVDF 膜可以重复使用。Y. 在做Western Blot时, PVDF膜用甲醇浸泡的目的? 解答: PVDF

传统WB面临的挑战



重复性差

1

➤ 5ug-20ug 总蛋白上样量

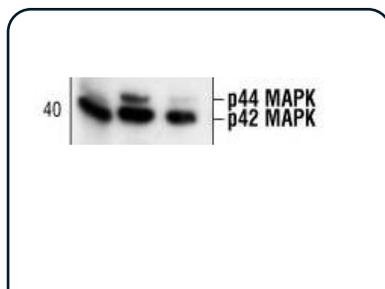
需要大量样本

2



大, 小分子量蛋白

3



定量差

4

➤ 12-24 小时, 手动操作多

周期长

5

传统WB面临的挑战



一直重复做WB

*图片来源于网络

MEET SIMPLE WESTERN

JESS ABBY WES



GEL-RUNNING AUTO
TRANSFER-FREE
BLOT-FREE
HANDS-FREE

超微量样品+自动化+定量

- 1 Simple Western 工作原理
- 2 Simple Western 优势及应用
- 3 Simple Western 实验操作简介

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SIMPLE WESTERN 工作原理

JESS ABBY WES



SIMPLE WESTERN 工作原理

化学发光检测

基于 CE-SDS 的免疫学分析



蛋白上样



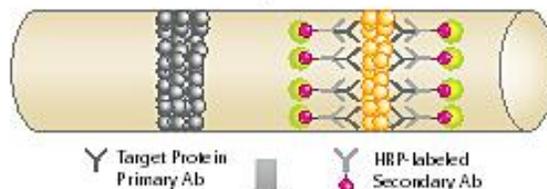
蛋白分离



蛋白固定



免疫杂交



数据采集



所有在毛细管内的步骤都
自动完成!

与传统 Western 不同:

无需制胶 (液态胶)

无需转膜 (紫外激活交联)

无需人工孵育/清洗/压片

无需人工数据分析

MEET SIMPLE WESTERN



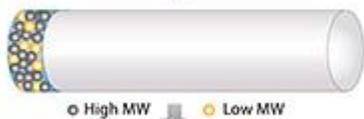
仪器型号	Wes	Abby	Jess
化学发光检测	√	√	√
荧光检测 (IR/NIR)	-	-	√
总蛋白检测	√	√	√
总蛋白归一化	-	-	√
Replex	-	√	√
化学发光成像	-	-	√

工作原理——荧光/总蛋白归一化

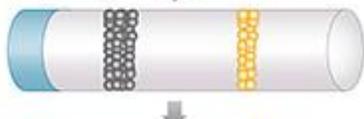
制胶



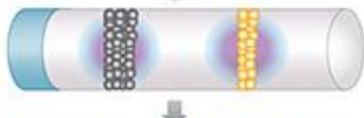
上样



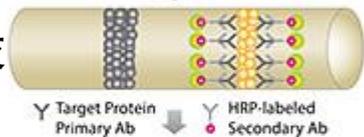
分离



固定



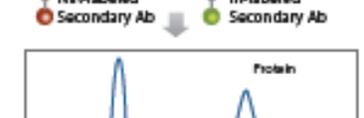
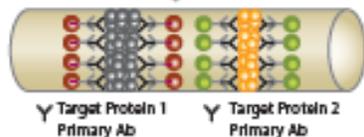
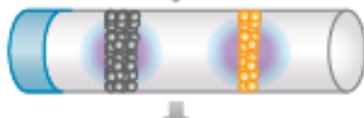
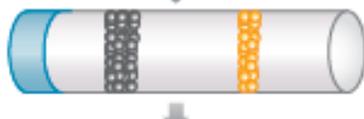
免疫杂交



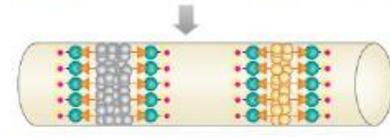
定量



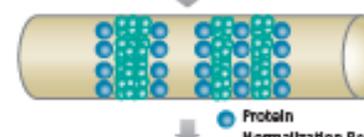
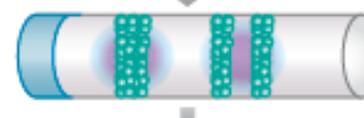
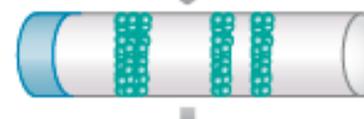
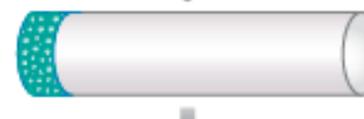
化学发光检测



荧光

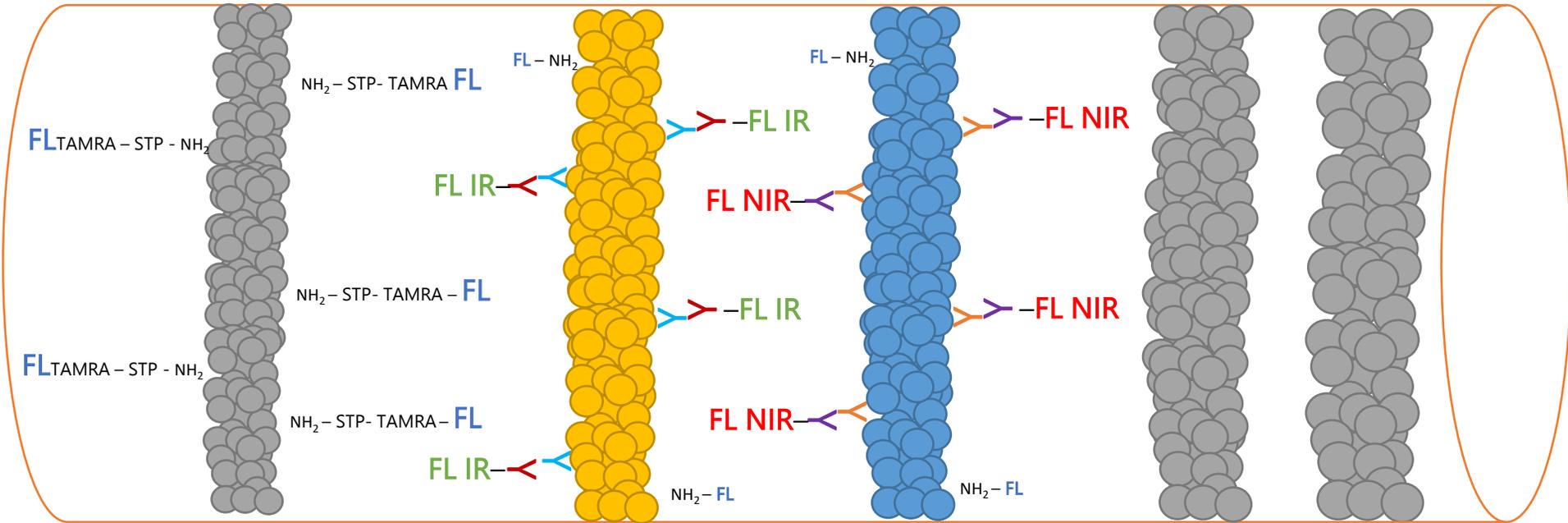


总蛋白检测



总蛋白归一化

总蛋白归一化检测



- 在 同一根 毛细管内
 - 基于荧光检测（与IR和NIR的荧光染料不同）
 - 可与所有蛋白质结合的专利试剂
 - 只需加入一种试剂
- 可检测毛细管结合的总蛋白量
- 校正**批次内**不同毛细管之间蛋白上样量的差异
- 必须和免疫检测同时使用

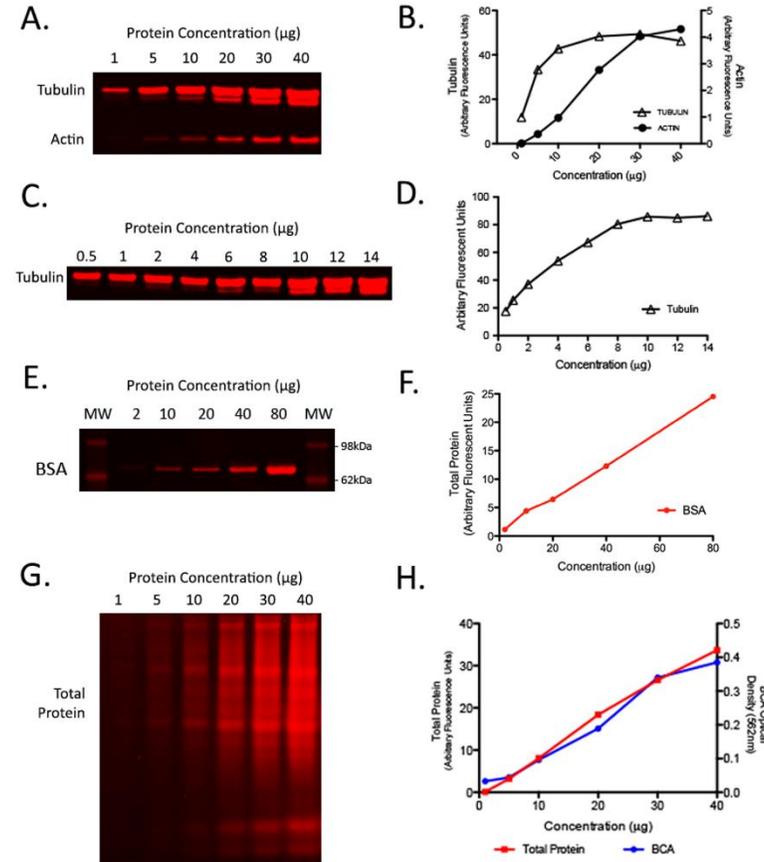
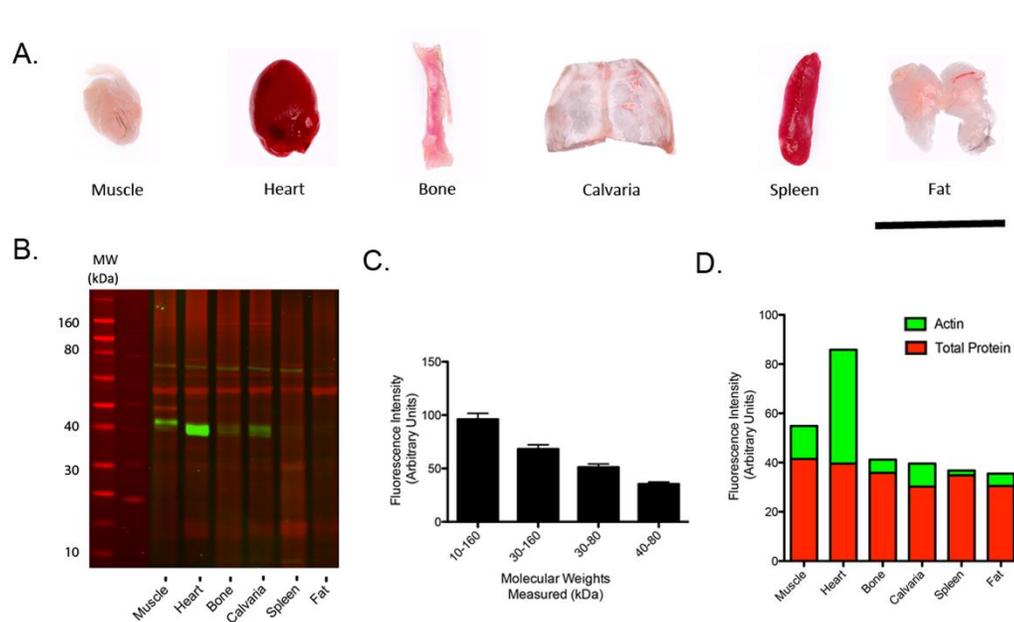
PROTEIN MOLECULAR WEIGHT RANGE		
LYSATE CONCENTRATION	12-230 kDa	
	Stock Solution	Reconstitution Agent
0.2-1.2 mg/mL	50 μ L	250 μ L

注意：-只能分析12-230kd的蛋白
 -上样浓度0.2-1.2mg/ml
 -不能单独检测

总蛋白归一化-WESTERN BLOT相对定量的趋势

内参蛋白受组织来源和实验处理的影响

内参蛋白的线性范围窄，无法满足所有靶蛋白的线性范围



Total Protein Analysis as a Reliable Loading Control for Quantitative Fluorescent Western Blotting, 2013. doi:10.1371/journal.pone.0072457.

总蛋白归一化的应用实例

PNAS

Proceedings of the
National Academy of Sciences
of the United States of America

RESEARCH ARTICLE

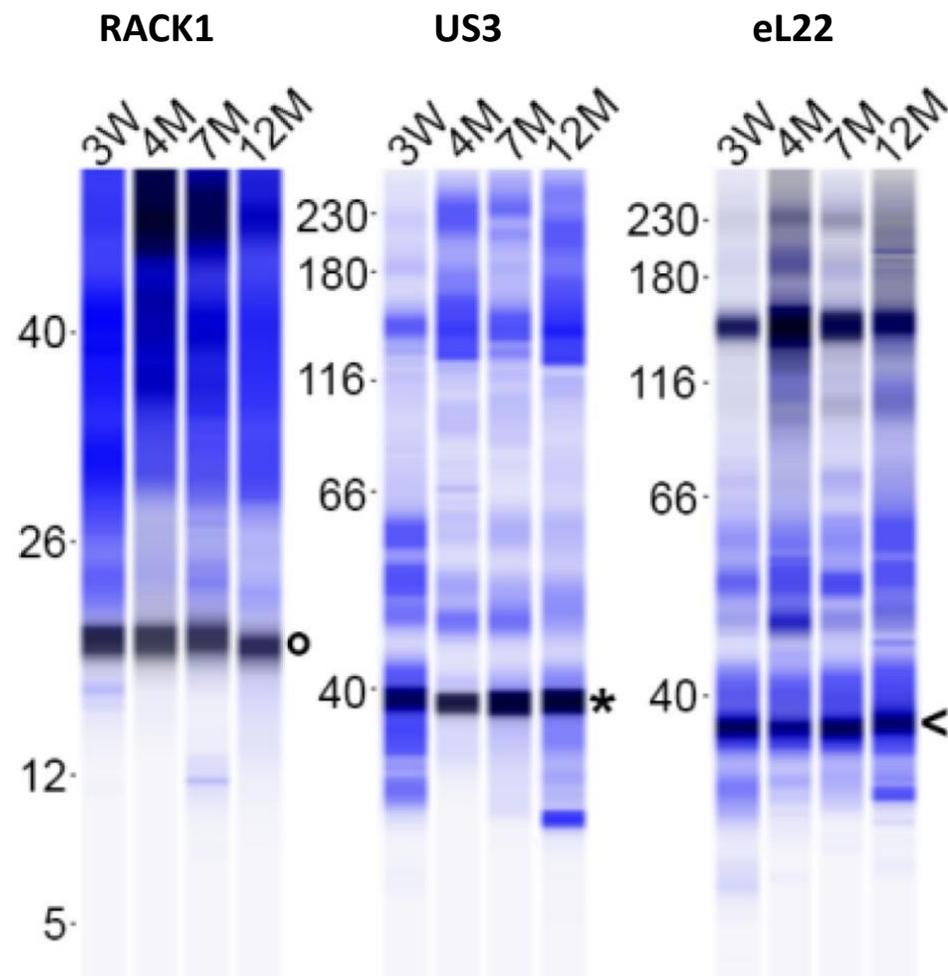
Invariable stoichiometry of ribosomal proteins in mouse brain tissues with aging

Susan Amirbeigi Arab, Parnian Kiani, Ana Velazquez Sanchez, Christoph Krisp, Andriy Kazantsev, Lars Fester, Hartmut Schlüter, and  Zoya Ignatova

PNAS November 5, 2019 116 (45) 22567-22572; first published October 21, 2019

<https://doi.org/10.1073/pnas.1912060116>

利用Jess的“总蛋白归一化”作为loading control，减少了不同样品间内参不平带来的相对定量误差。



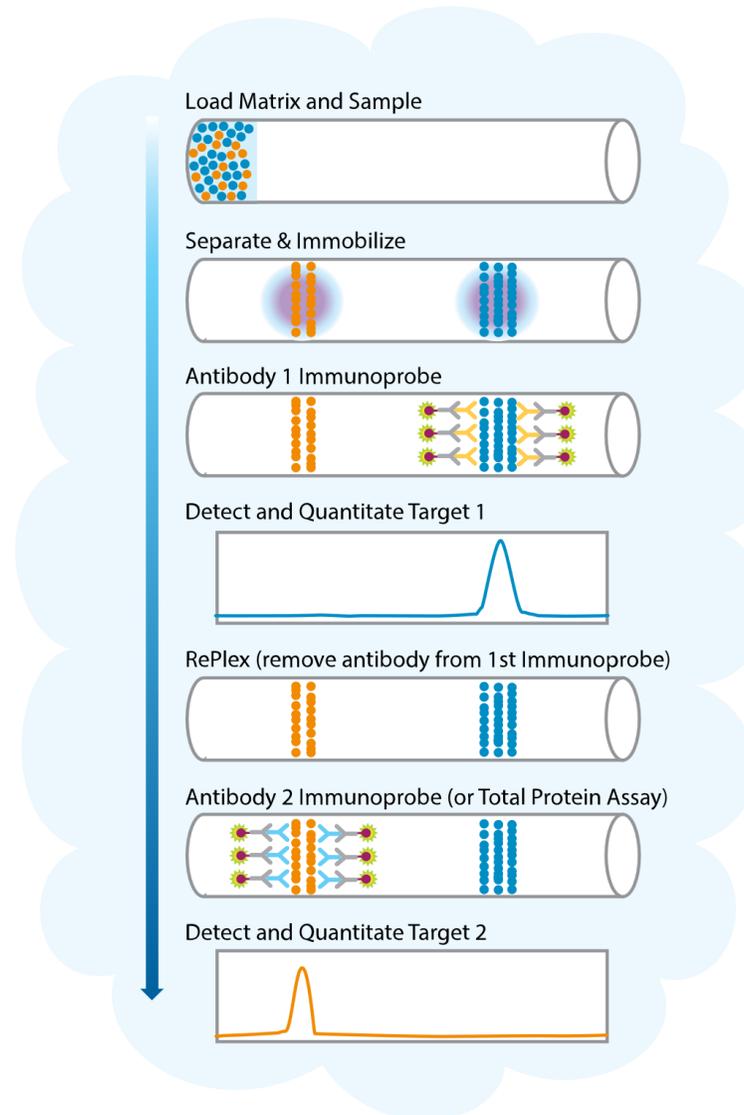
RePlex 定义

一根毛细管内进行两轮免疫学检测 (2 次连续检测, 每轮只测1个靶点)

可用于化学发光、荧光及总蛋白检测

节省单个数据点检测费用

利用Simple Western技术独有的蛋白质与毛细管共价交联固定



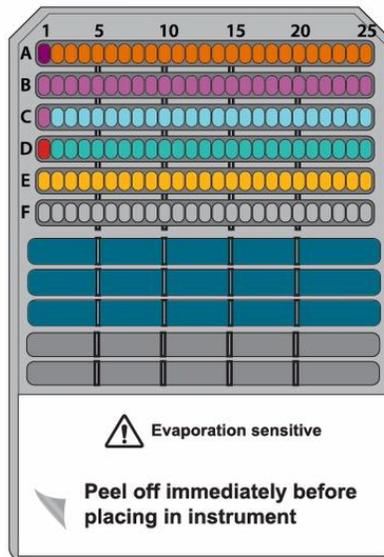
您需要做的工作



1. 配制样品和试剂

- 加样
- 混匀
- 加热

*样品不能加 Loading Buffer



2. 加样

- 加marker 
- 加样品 
- 加封闭液 
- 加一抗 
- 加二抗 
- 加发光液 
- 加洗液 



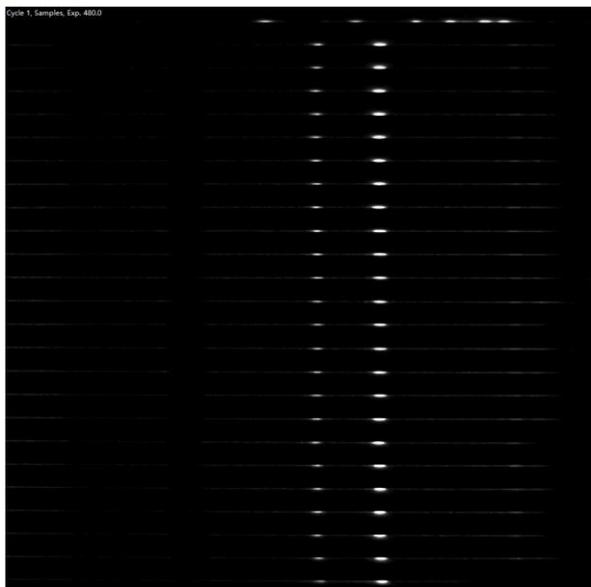
3. 运行

- 放毛细管卡盒
- 放加样板
- 运行程序

THREE HOURS

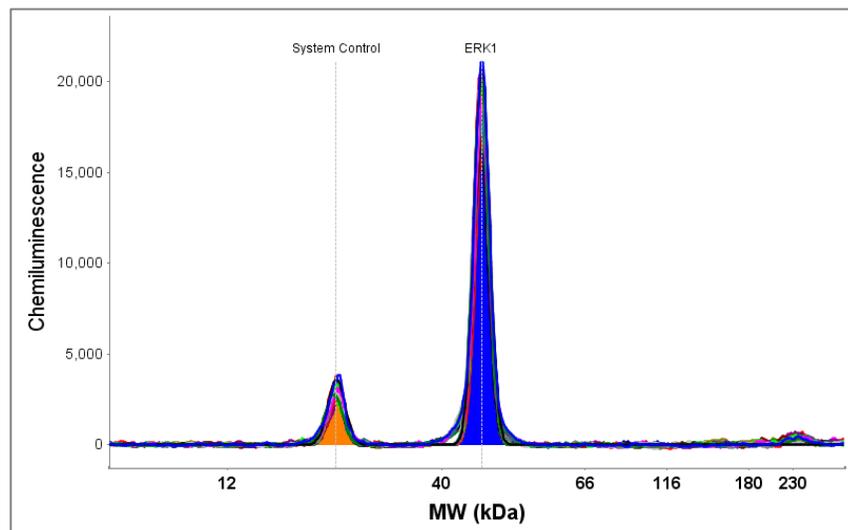
LATER...

三种结果展现形式



成像图

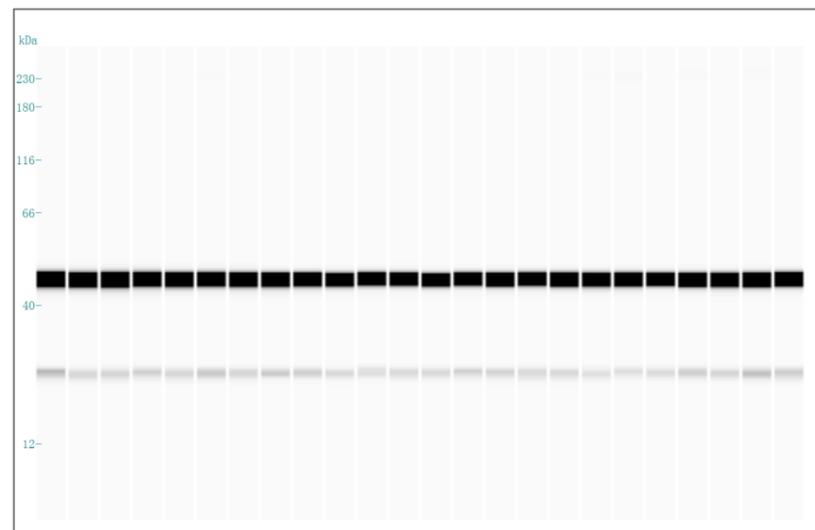
毛细管实际曝光图



峰形图

峰面积为
化学发光信号值

第一次实现直接获得条带化学发光信号值



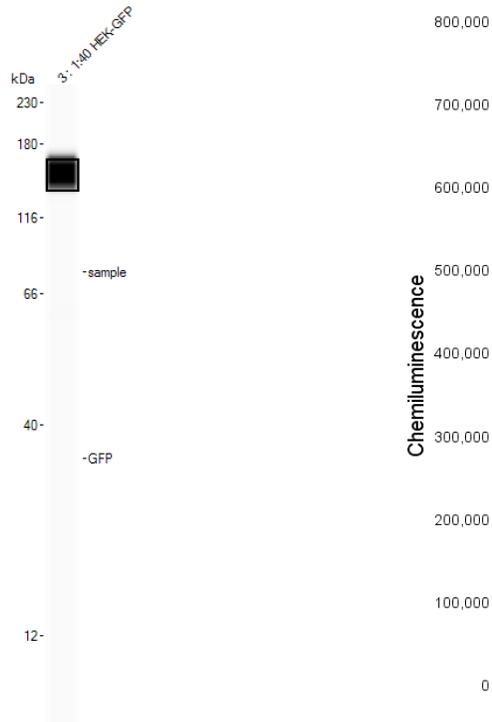
泳道图

条带模拟图

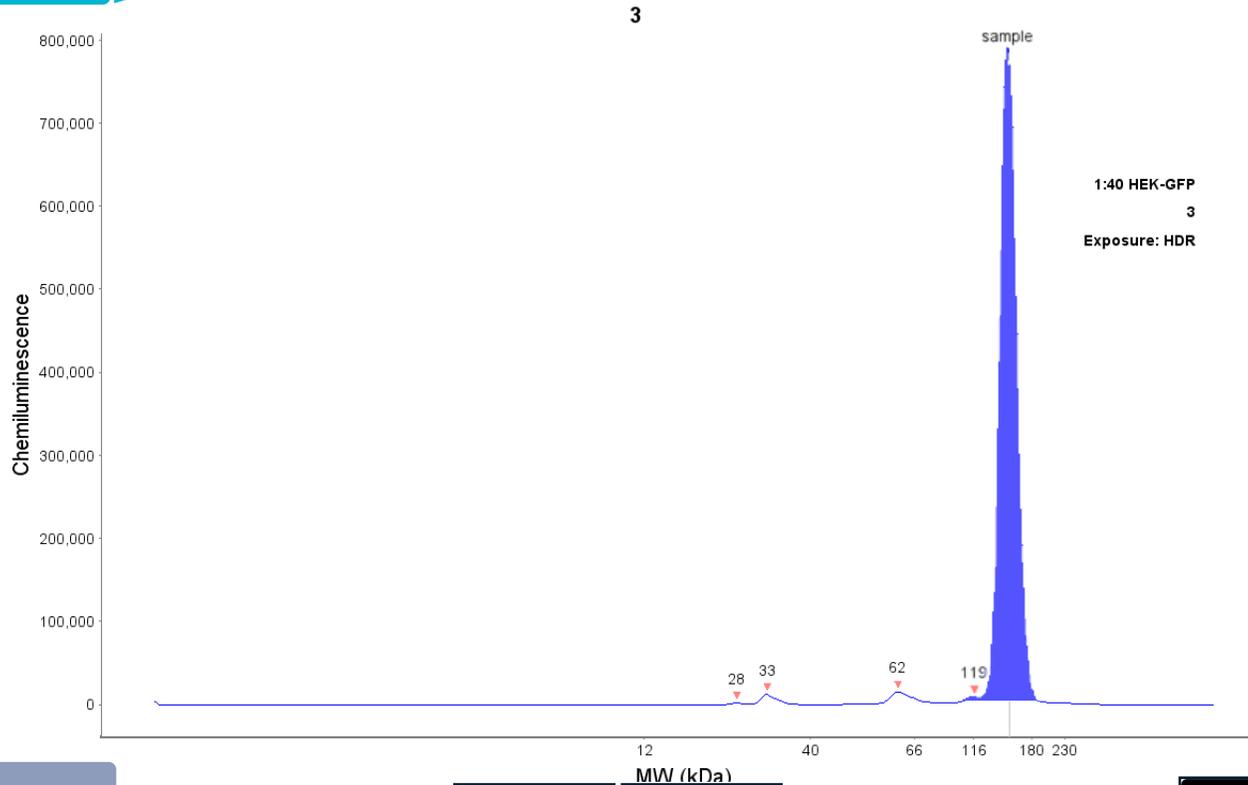
数字化WESTERN

1. Lane View

More Detail



2. Electropherogram View



3. More Rich Information

Sample	Primary	Cap	Peak	Name	Position	MW (kDa)	Peak Height	Peak Area	% Area	Conc (ng...)	Width	Peak Area S/N	Baseline
							Height	Area				S/N	Baseline
1:40 HEK-GFP	Gt anti...	3	1		422	28	2386.9	25194			9.9	84.5	308.4
1:40 HEK-GFP	Gt anti...	3	2		445	33	11712.6	170768			13.7	392.8	322.4
1:40 HEK-GFP	Gt anti...	3	3		540	62	16282.4	276943			16.0	422.1	386.7
1:40 HEK-GFP	Gt anti...	3	4		594	119	12403.2	178493			13.5	120.6	419.4
1:40 HEK-GFP	Gt anti...	3	5	sample	620	155	763025.7	9340434	100.0		11.5	29468.0	432.1

MEET SIMPLE WESTERN

JESS ABBY WES

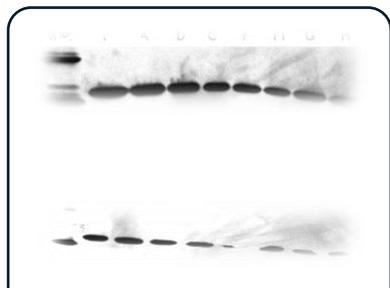


GEL-RUNNING AUTO
TRANSFER-FREE
BLOT-FREE
HANDS-FREE

超微量样品+自动化+定量

- 1 Simple Western 工作原理
- 2 Simple Western 优势及应用
- 3 Simple Western 实验操作简介

传统WB面临的挑战



重复性差

1

➤ 5ug-20ug 总蛋白上样量

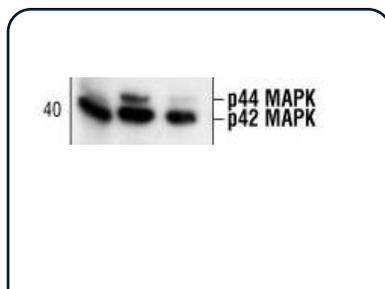
需要大量样本

2



大, 小分子量蛋白

3



定量差

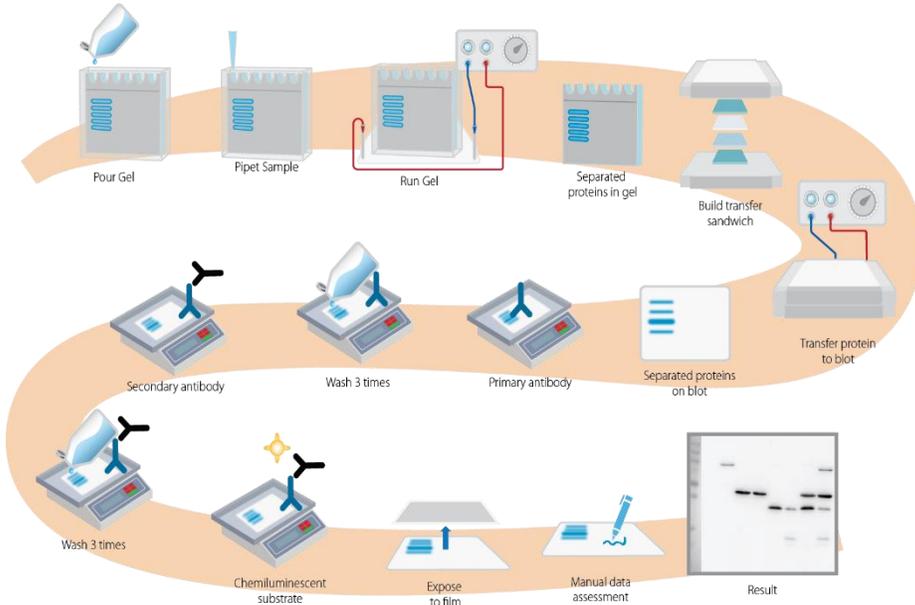
4

➤ 12-24 小时, 手动操作多

周期长

5

1. SIMPLE WESTERN – 重现性好



The error in quantitative gel electrophoresis/Western blotting was investigated considering the purity testing of erythropoietin. **The overall error was over 35% relative standard deviation.** However, an analysis of variance elucidated that the interoperator variability was the dominant error source, which already explained almost 80% of the total variance. Careful compilation and investigation of the possible error sources strongly indicates that the immunoreaction after blotting and the subsequent color reaction are the major error sources in this case.

Roche CV > 35%

Koller, A. and H. Watzig (2005). "Precision and variance components in quantitative gel electrophoresis." *Electrophoresis* **26**(12): 2470-2475.

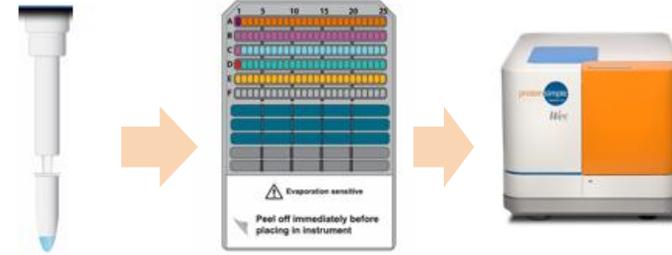


Table 1. Reproducibility (intermediate precision) of Simon™

Experiment	Concentration in $\mu\text{g/mL}$			
	Sample 1	Sample 2	Sample 3	
			p106	p86
Run 1	100	219	138	60
Run 2	102	256	138	53
Run 3	106	220	125	63
Run 4	111	242	131	53
Run 5	94	238	121	60
Mean	103	235	130	57
SD	7	16	8	5
%CV	6	7	6	9

MERCK CV = 9%

Rustandi, R. R., et al(2012). "Qualitative and quantitative evaluation of Simon, a new CE-based automated Western blot system as applied to vaccine development." *Electrophoresis* **33**(17): 2790-2797.

2. SIMPLE WESTERN – 微量样品优势—分选干细胞

应用Wes完成2000个分选骨髓干细胞western blot

The impact of integrin $\beta 2$ on granulocyte/macrophage progenitor proliferation

LI-JIE ZHANG,^a CEN YAN,^a SARAH SCHOUTEDEN,^b XIAO-JUAN MA,^a DONG ZHAO,^a THORSTEN PETERS,^c CATHERINE M. VERFAILLIE,^b YING-MEI FENG^{ib,a,b}

Key Words. granulocyte/macrophage progenitors • integrin $\beta 2$ • Fc ϵ R1 α • GATA2 • proliferation

传统Western Blot检测:

一只小鼠流式分选获取骨髓干细胞量: 1000个左右

1个孔需要数以万记细胞裂解的总蛋白, 需要数十只小鼠, 代价高昂。

Wes检测: 1000个细胞 (相等1只小鼠) 即可检测到GAPDH信号, 2500个细胞 (相等于2-3小鼠) 即可获得比较强且稳定的信号。

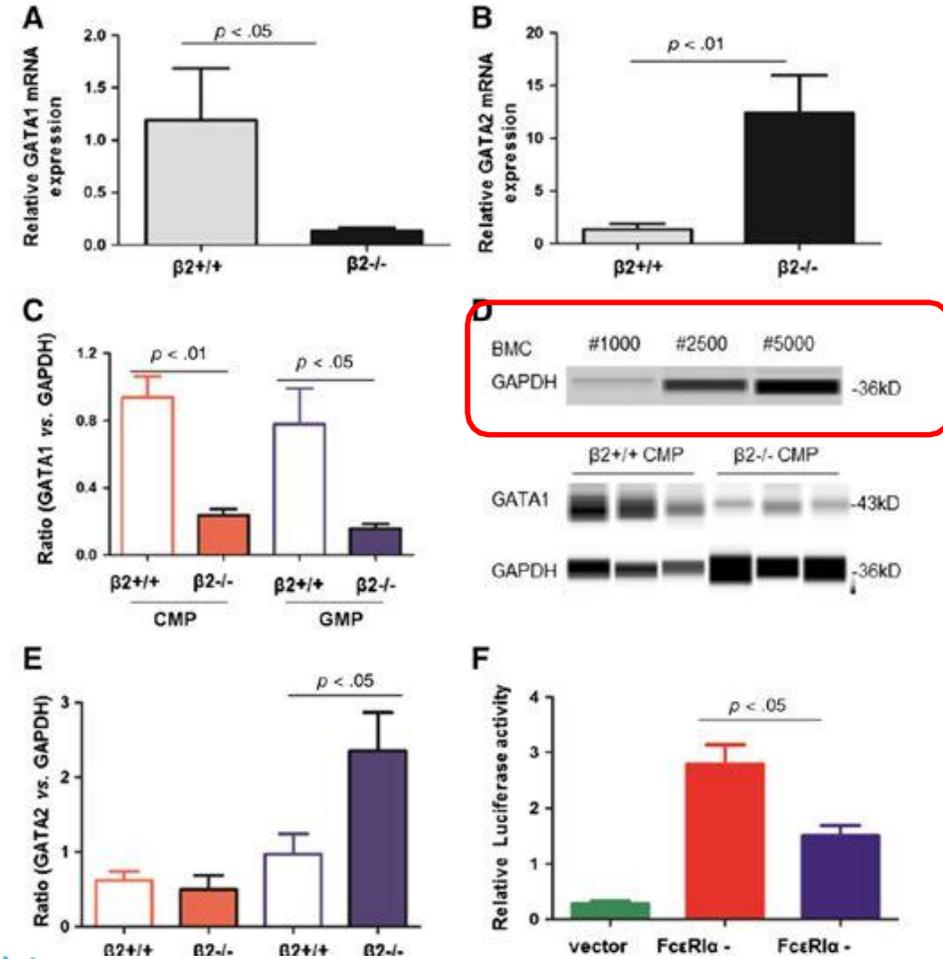
费用只有传统方法几分之一

Transcriptional regulation of Fc ϵ R1 α by GATA2. GMP were sorted out by FACS.

After protein concentration determination, equal amount of proteins was separated by capillary western blot.

GATA1, GATA2 and GAPDH expression were studied. Each sample was pooled from 2 to 4 mice.

The impact of integrin $\beta 2$ on granulocyte/macrophage progenitor proliferation. **STEM CELLS 2018**



首都医科大学附属潞河医院内分泌中心
糖尿病防治北京市重点实验室

2. SIMPLE WESTERN – 微量样品优势—活检样本

nature

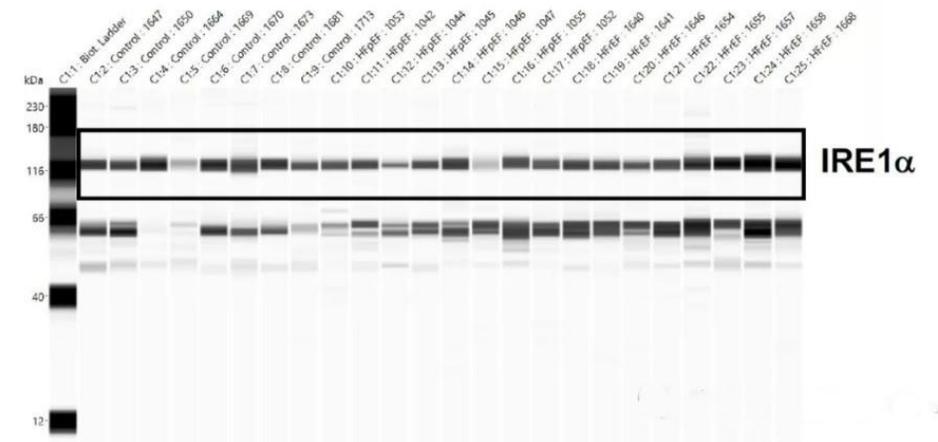
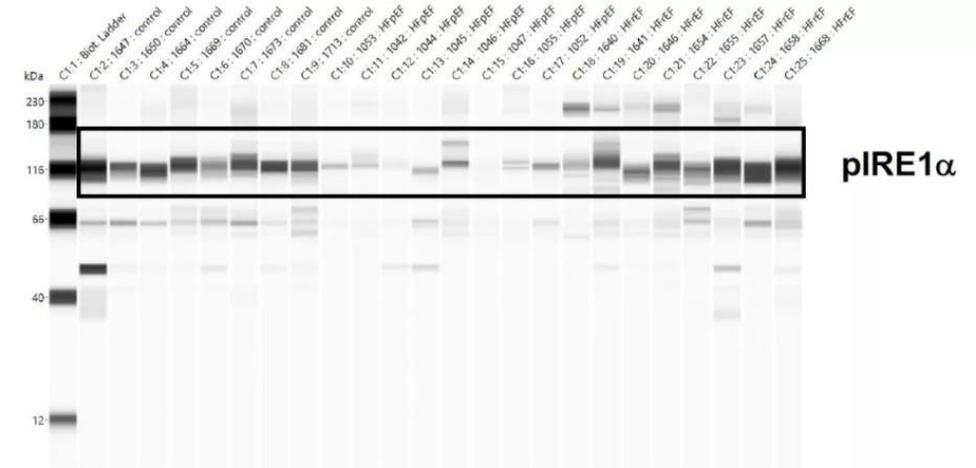
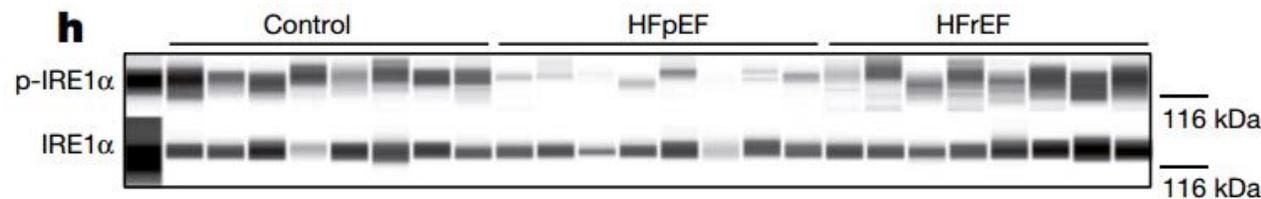
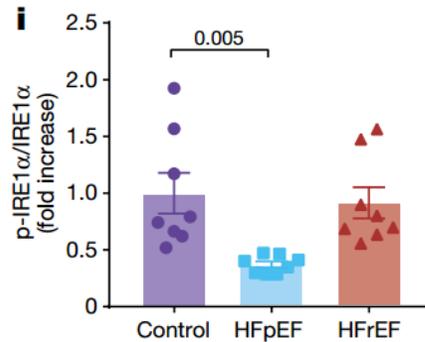
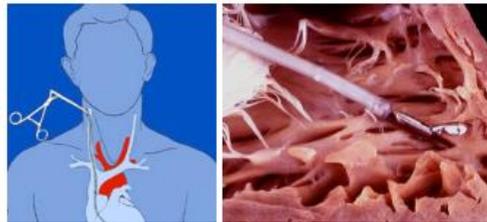
Article | Published: 10 April 2019

Nitrosative stress drives heart failure with preserved ejection fraction

Gabriele G. Schiattarella^{1,2}, Francisco Altamirano¹, Dan Tong¹, Kristin M. French¹, Elisa Villalobos¹, Soo Young Kim¹, Xiang Luo¹, Nan Jiang¹, Herman I. May¹, Zhao V. Wang¹, Theodore M. Hill¹, Pradeep P. A. Mammen¹, Jian Huang¹, Dong I. Lee³, Virginia S. Hahn³, Kavita Sharma³, David A. Kass³, Sergio Lavandero^{1,4,5}, Thomas G. Gillette¹ & Joseph A. Hill^{1,6*}

Wes检测人心脏活检样本 IRE1 α - XBP1 信号通路蛋白

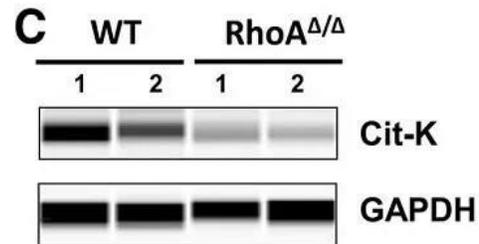
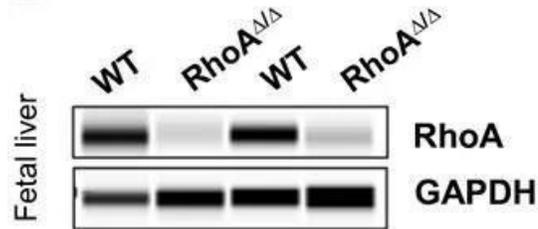
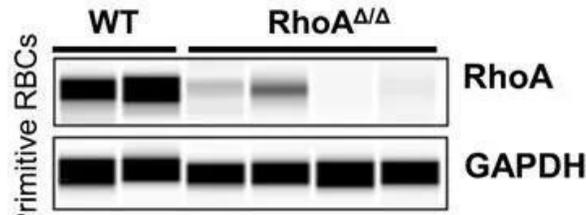
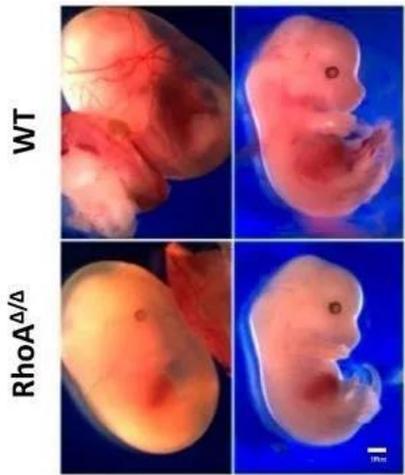
右心室心内膜活检术



全膜western blot结果

2. SIMPLE WESTERN – 微量样品优势—胚胎来源样本

小鼠胚胎分选红细胞中分裂关键蛋白检测

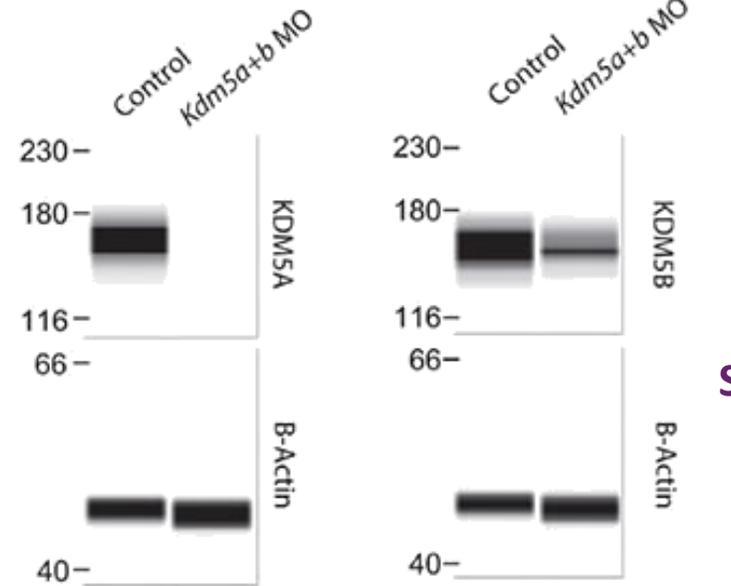
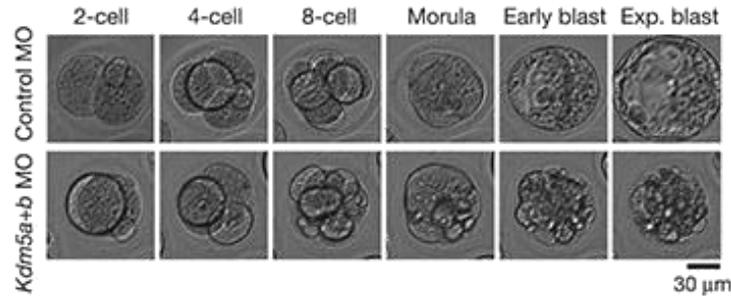


美国辛辛那提大学辛辛那提儿童医学中心

Cytokinesis failure in RhoA-deficient mouse erythroblasts involves actomyosin and midbody dysregulation and triggers p53 activation.

BLOOD

胚胎发育研究中组蛋白检测



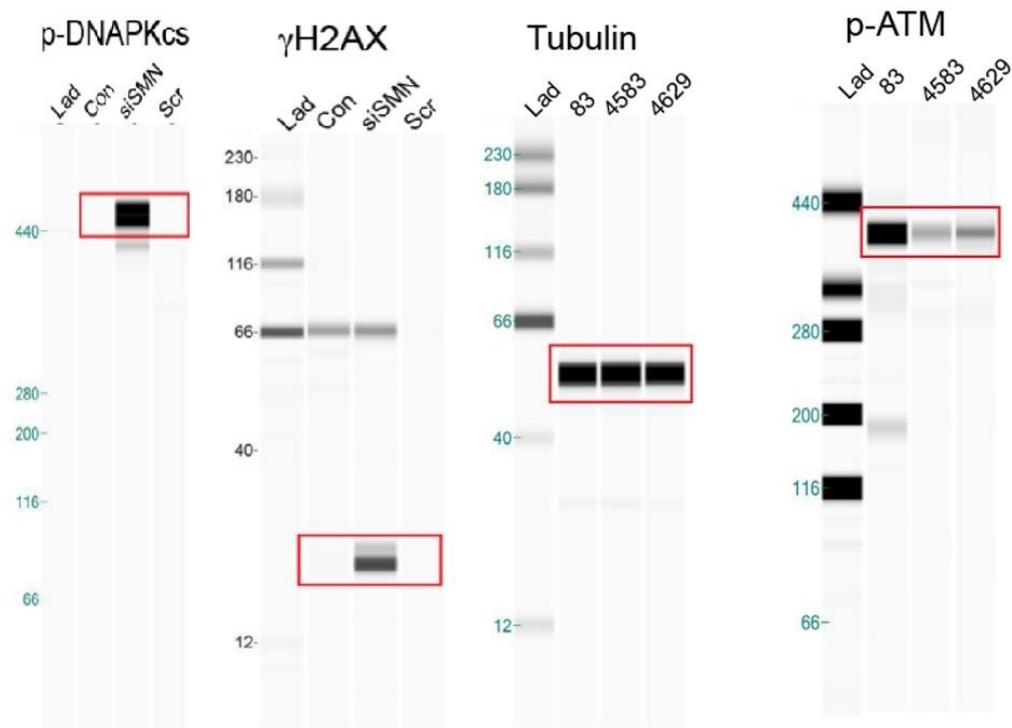
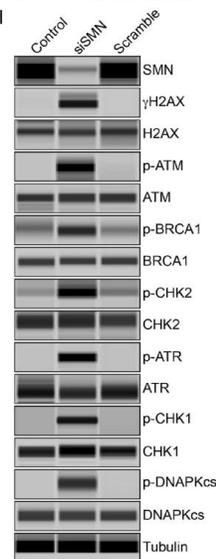
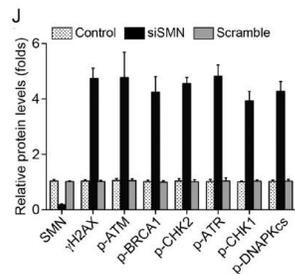
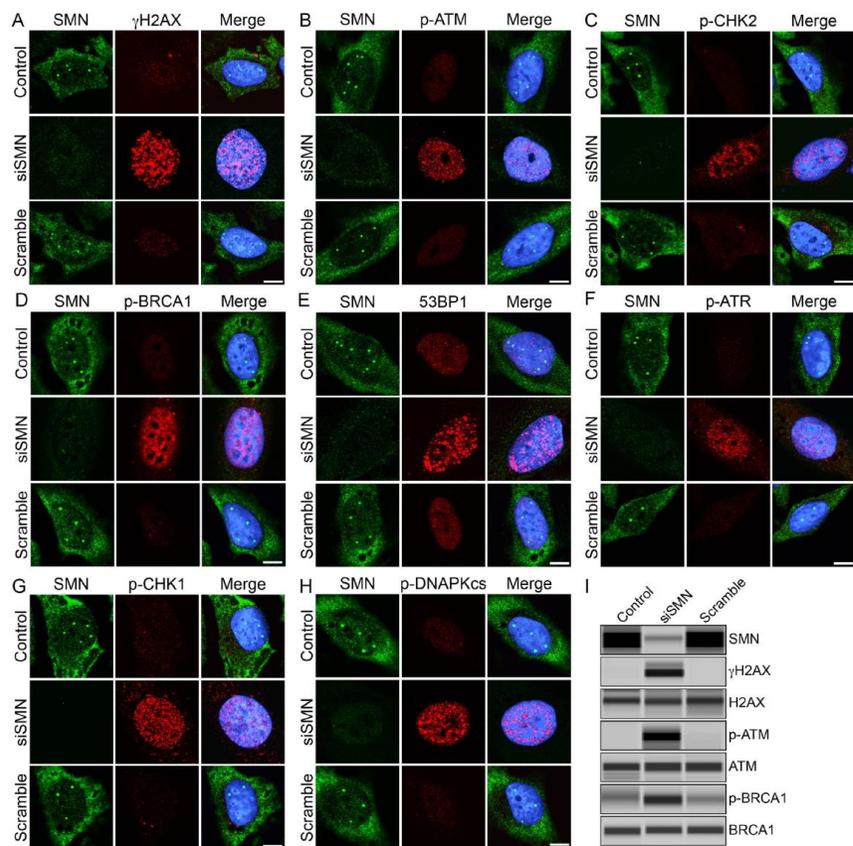
Broad histone H3K4me3 domains in mouse oocytes modulate maternal-to-zygotic-transition, **Nature**.(2016)

nature

挪威奥斯陆大学附属医院
+
美国加州路德维格癌症研究所

132个2-cell胚胎中，
得到7 ul的样品
Simple wes系统：上样3 ul，
实际消耗40 nl。

3. SIMPLE WESTERN –蛋白质研究不再纠结分子量大小



脊髓性肌萎缩 (简称SMA疾病) 相关蛋白:

SMN (40KD) **DNAPKcs (450KD)** **CHK1 (56KD)** **ATR (300KD)** **CHK2 (62KD)** **BRCA1 (220KD)** **ATM (350KD)** **H2Ax (15KD)** **rH2AX (15KD)** **SETX (35KD)** **Tubulin (55KD)**

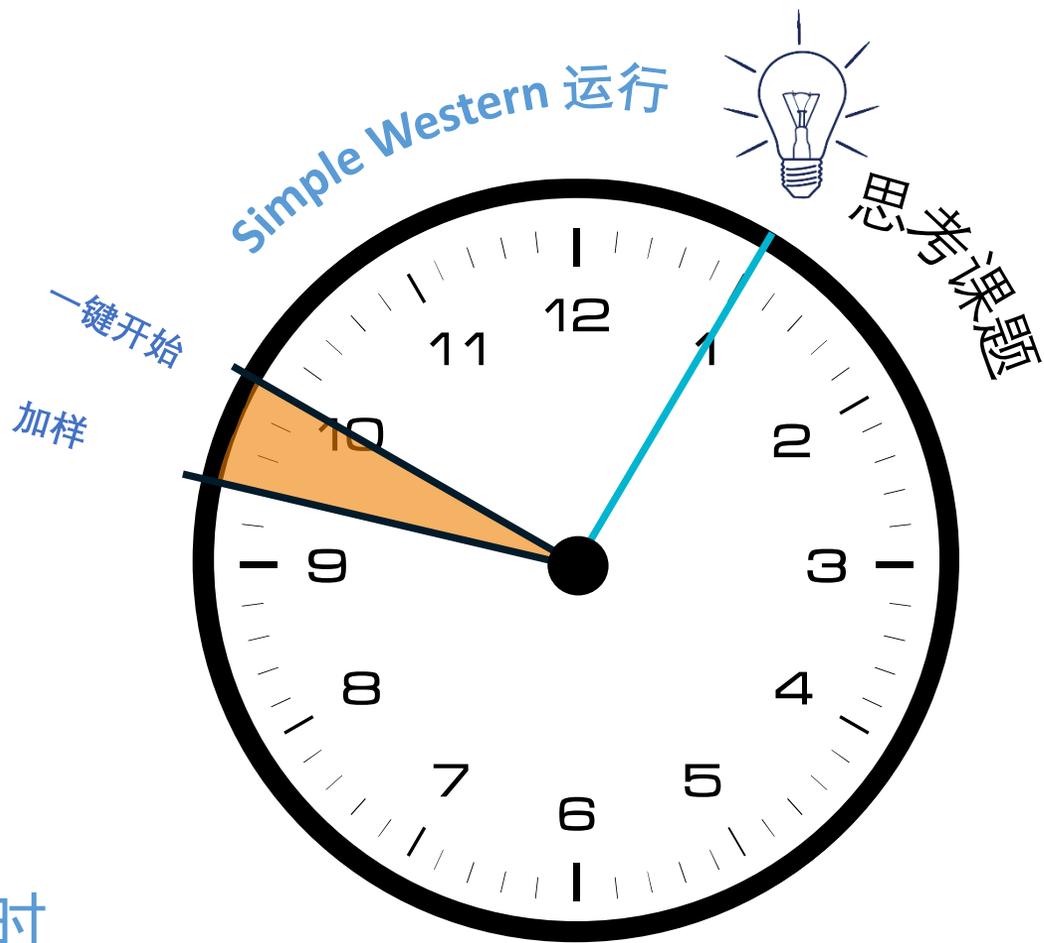
Combined deficiency of Senataxin and DNA-PKcs causes DNA damage accumulation and neurodegeneration in spinal muscular atrophy. Nucleic Acids Research, 2018

4. SIMPLE WESTERN – 3 H 完成实验

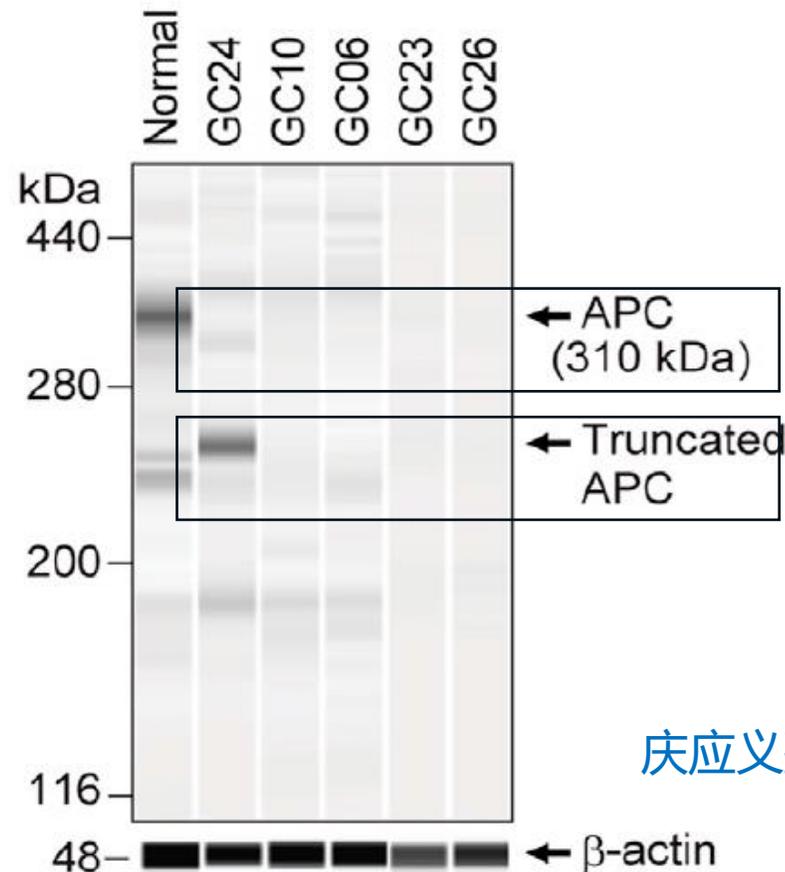
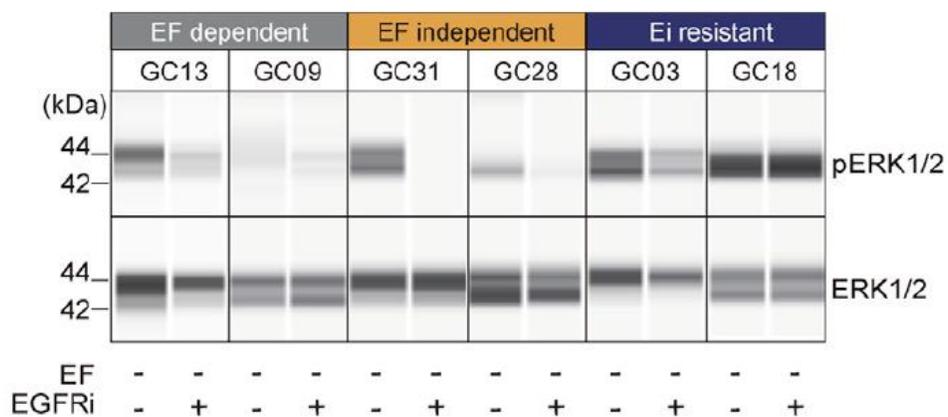
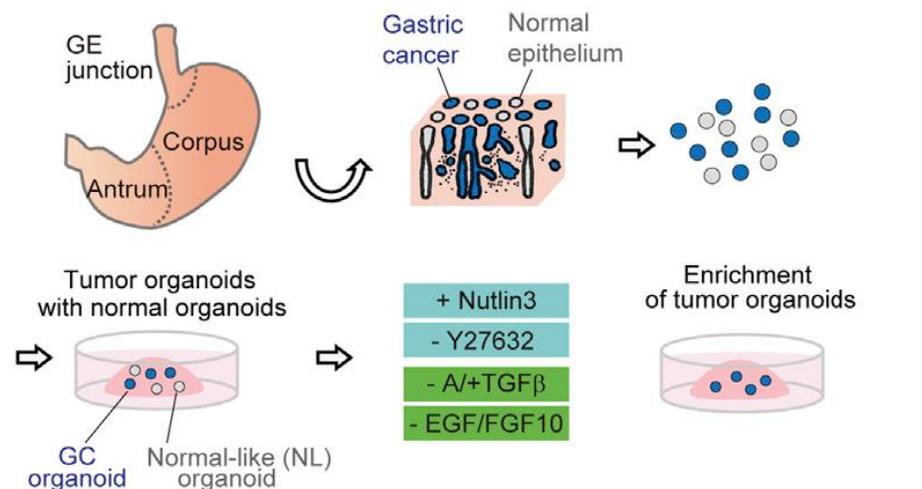


25 个样品
3 个小时出结果

从开始运行到给出结果仅需 **3** 小时
人工操作时间仅 **0.5** 小时



5. SIMPLE WESTERN – 内参蛋白相对定量



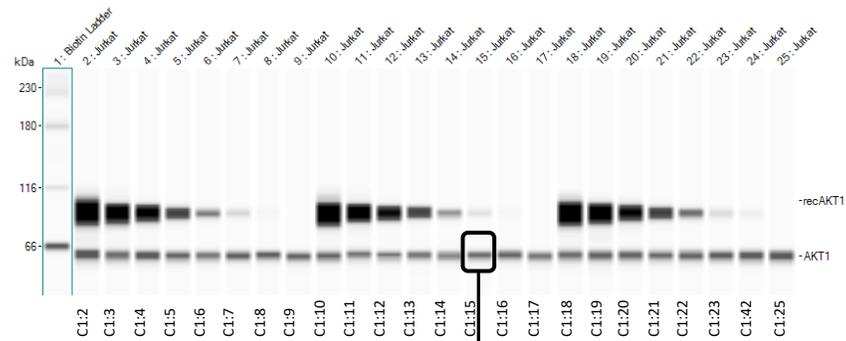
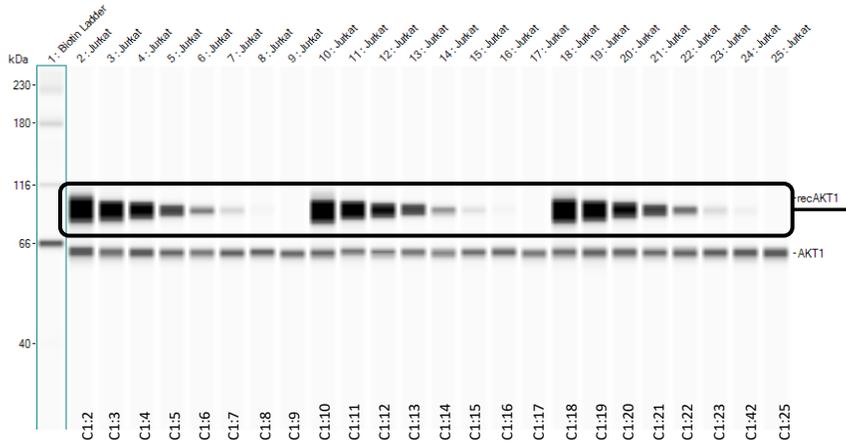
胃癌类器官研究
庆应义塾大学医学院消化内科

Divergent Routes toward Wnt and R-spondin Niche Independence during Human Gastric Carcinogenesis.
Nanki et al., 2018, *Cell* 174, 856–869 August 9, 2018



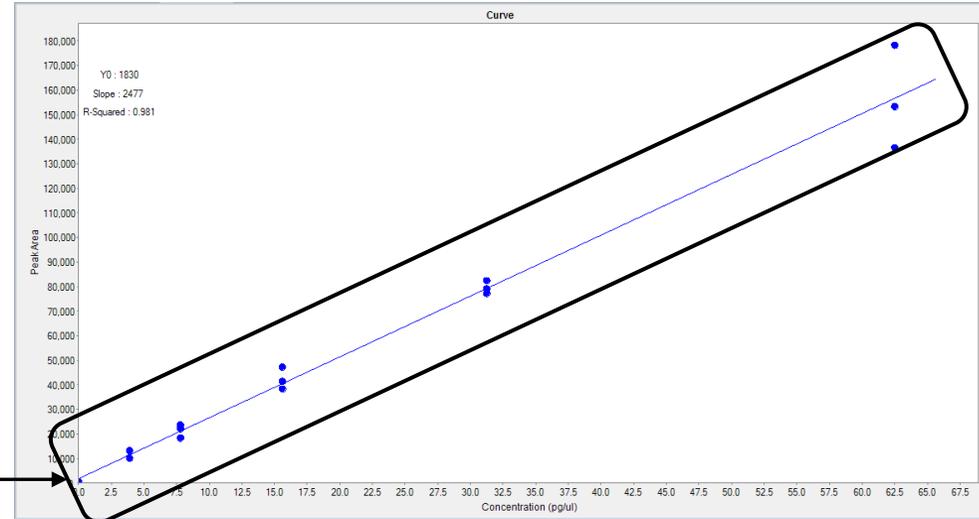
5. SIMPLE WESTERN – 绝对定量

1. Confirm with Lane View



GST标签的重组AKT蛋白

2. Create Standard Curve



3. Quantitate

Sample	Prim...	Cap	Peak	Name	Positi...	MW (...)	Height	Area	% Area	Conc ...	Width	S/N	Baseline
Jurkat	Anti...	C1:2	1	AKT1	496	64	4874.3	70846	14.4	27.86	13.7	23...	129.4
Jurkat	Anti...	C1:2	2	rec...	533	96	2980...	4209...	85.6	169.2	13.3	13...	142.6
Jurkat	Anti...	C1:3	1	AKT1	495	64	4193.2	58858	18.5	23.02	13.2	30...	132.9
Jurkat	Anti...	C1:3	2	rec...	533	95	2182...	2596...	81.5	104.1	11.2	16...	135.3
Jurkat	Anti...	C1:4	1	AKT1	495	64	4912.3	57228	24.3	22.37	10.9	23...	97.4
Jurkat	Anti...	C1:12	2	rec...	535	94	1320...	1365...	75.7	62.50	9.7	67...	123.3
Jurkat	Anti...	C1:13	1	AKT1	496	64	4235.3	44507	36.6	17.23	9.9	30...	113.4
Jurkat	Anti...	C1:13	2	rec...	534	95	7360.9	77229	63.4	31.25	9.9	52...	104.6
Jurkat	Anti...	C1:14	1	AKT1	498	63	3970.7	44205	53.6	17.11	10.5	30...	129.2
Jurkat	Anti...	C1:14	2	rec...	535	94	3681.9	38299	46.4	15.60	9.8	27...	129.9
Jurkat	Anti...	C1:15	1	AKT1	499	64	4238.4	45984	71.5	17.83	10.2	30...	107.4

17.83 (pg/ul)

5. SIMPLE WESTERN – 总蛋白均一化

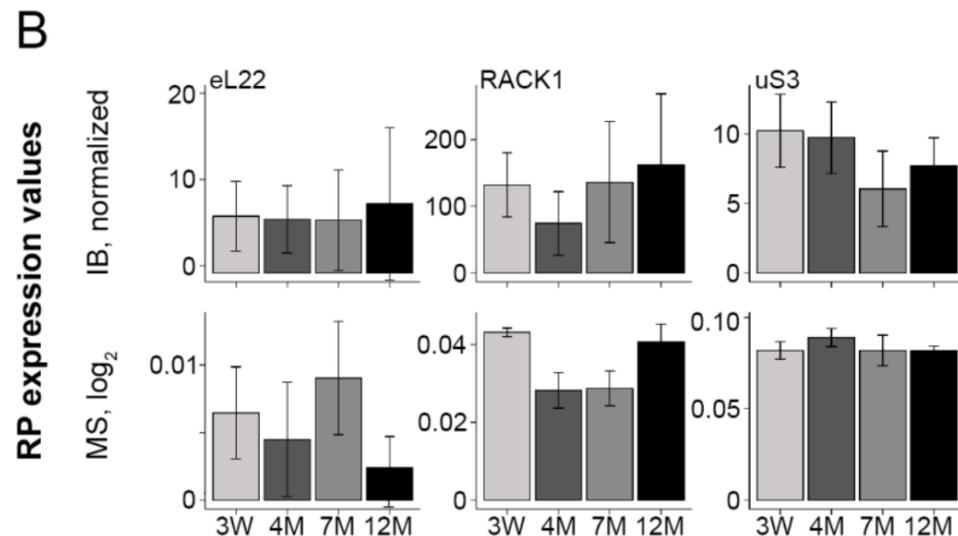
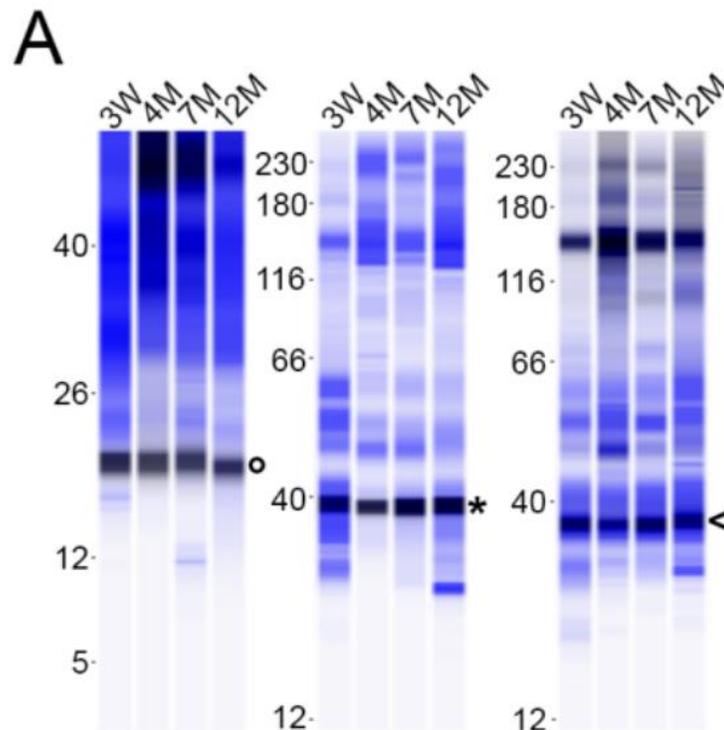
PNAS

Proceedings of the
National Academy of Sciences
of the United States of America

➤ PNAS: 使用Jess总蛋白归一化技术定量研究核糖体蛋白

德国汉堡大学
生物化学与分子生物学系

本研究利用Jess, 对不同年龄的小鼠肝脏组织中的核糖体蛋白 (RPs) 进行定量, 进一步从蛋白水平上, 揭示了无论是不同年龄的小鼠, 还是小鼠不同组织 (脑组织亦或是肝脏组织), 其核糖体蛋白的蛋白水平均保持不变。



Immunoblot Analysis. The immunoblot analysis was performed with liver samples using the capillary electrophoresis system (Jess, ProteinSimple). Frozen tissues ($n = 4$ to 6 biological replicates) were homogenized separately and fractionated in monosomal and polysomal fractions as prepared for the mass spectrometry analysis (see above) and loaded on the Jess separation module (2 to 40 kDa and 12 to 230 kDa). Antibodies against RACK1, uS3, and eL22 were purchased from Santa Cruz Biotechnology. The corresponding RP peaks were normalized to the total protein concentration using a protein normalization kit (ProteinSimple) and quantified using the Jess quantification module.

利用Jess的“总蛋白归一化”作为loading control, 减少了不同样品间内参不平带来的相对定量误差。

Invariable stoichiometry of ribosomal proteins in mouse brain tissues with aging. *Proc Natl Acad Sci.* 2019 Nov 5;116(45):22567-22572.

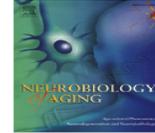
6. 全膜 WESTERN BLOT



Contents lists available at ScienceDirect

Neurobiology of Aging

journal homepage: www.elsevier.com/locate/neuaging

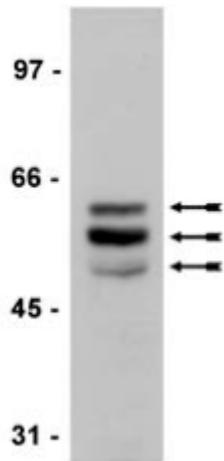


稳定的出现的条带迁移，可能是新发现

Sustained high levels of neuroprotective, high molecular weight, phosphorylated tau in the longest-lived rodent

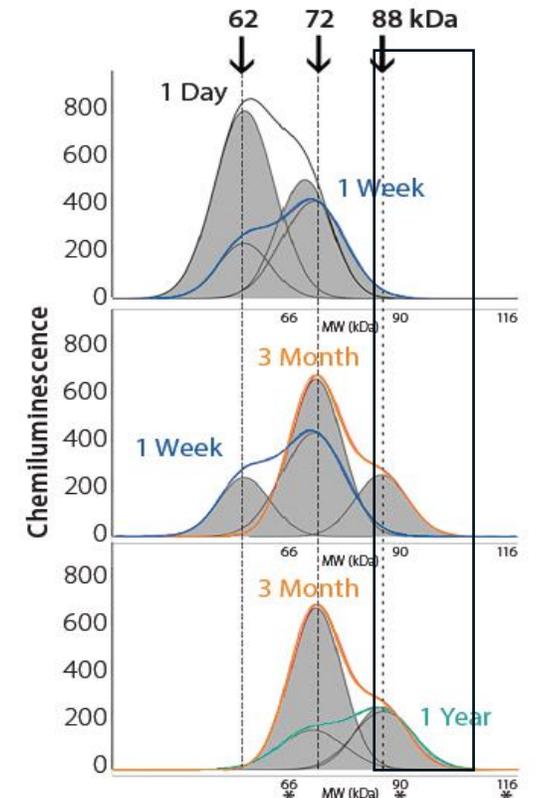
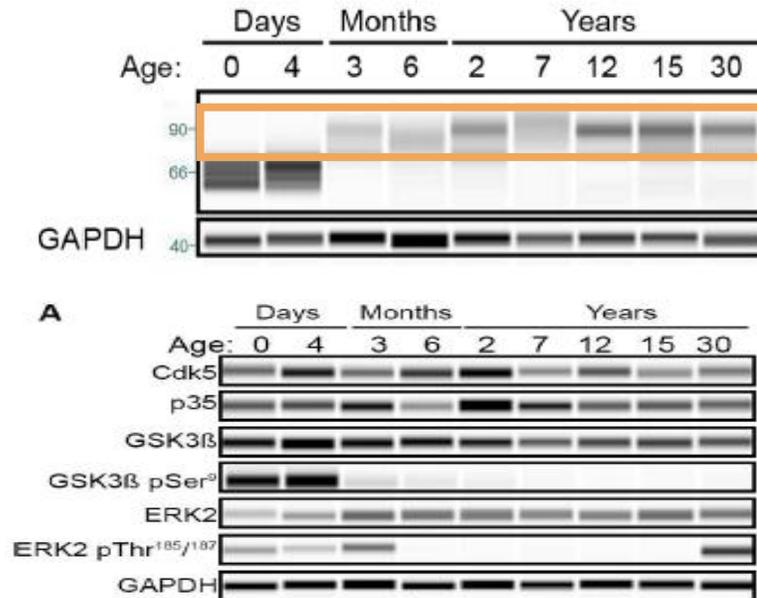
Miranda E. Orr, Valentina R. Garbarino, Angelica Salinas, Rochelle Buffenstein*

Department of Physiology and The Barshop Institute for Longevity and Aging Studies, University of Texas Health Science Center at San Antonio, San Antonio, TX, USA



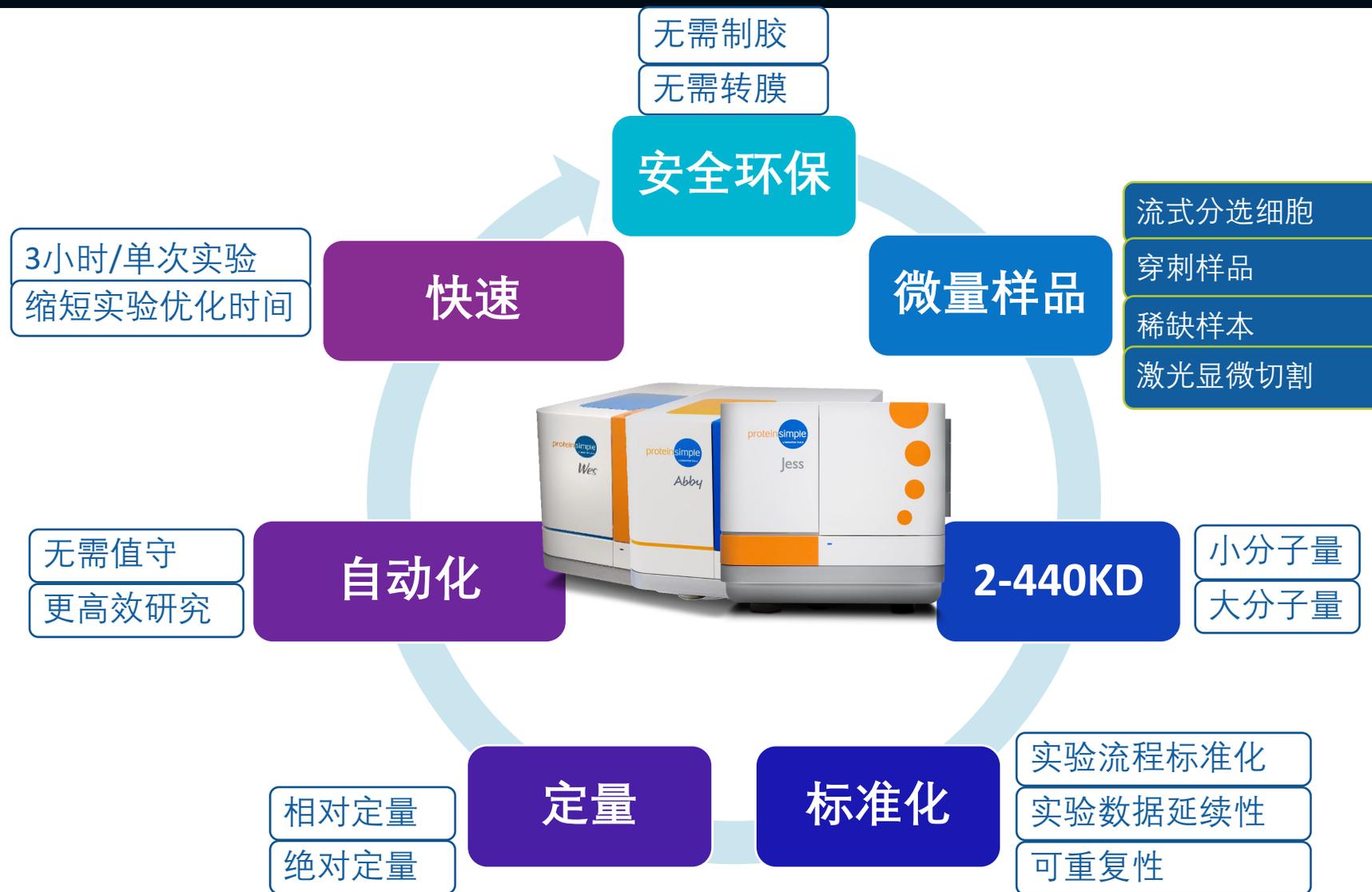
Specificity

Recognizes Tau (4-repeat isoform RD4), Mr 45-65 kDa. Higher MW band (68-72 kDa) represents phosphorylated Tau.



随着Rat 年龄增大，88KD蛋白出现，且在Tau蛋白中占的比例越来越高。

SIMPLE WESTERN 特点



MEET SIMPLE WESTERN

JESS ABBY WES



GEL-RUNNING AUTO
TRANSFER-FREE
BLOT-FREE
HANDS-FREE

超微量样品+自动化+定量

- 1 Simple Western 工作原理
- 2 Simple Western 优势及应用
- 3 Simple Western 实验操作简介

试剂盒

荧光检测试剂盒

Separation Module

- 2-40kDa分离试剂盒
- 12-230kDa分离试剂盒
- 66-440kDa 分离试剂盒
- 25 或 13 毛细管卡盒



Detection Module

- Anti-Rabbit NIR
- Anti-Mouse NIR
- Anti-Rabbit IR
- Anti-Mouse IR



总蛋白均一化试剂盒

Protein Normalization Assay Module for Jess (AM-PN01)



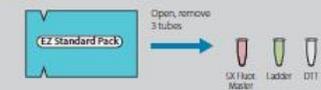
RePlex Kit – RP-001



操作说明书

1. Assay Module preparation

A PREPARE STANDARD PACK REAGENTS

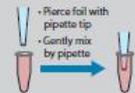


DTT (Clear Tube)



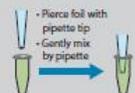
- Add 40 μ L deionized water to make a 400 mM solution

Fluorescent 5X Master Mix (Pink Tube)



- Add 20 μ L 10X Sample Buffer
- Add 20 μ L prepared 400 mM DTT solution

Biotinylated Ladder (Green Tube with Pink Pellet)



- Add 20 μ L deionized water

B PREPARE YOUR SAMPLES

- The optimal protein concentration depends on the expression level of your protein. Dilute lysate as necessary with 0.1X Sample Buffer.
- Combine 1 part 5X Fluorescent Master Mix with 4 parts diluted lysate in a microcentrifuge tube (final concentration 0.4 mg/mL for chemiluminescence or 1.0 mg/mL for fluorescence). Produce enough diluted sample volume required for assay.



C DENATURE YOUR SAMPLES



D MIX LUMINOL-S AND PEROXIDE (IF APPLICABLE)

- Combine 200 μ L Luminol-S and 200 μ L Peroxide in a microcentrifuge tube



E PREPARE PROTEIN NORMALIZATION REAGENTS

- The Protein Normalization Reagent should only be prepared immediately before loading the plate.



- Prepare the Protein Normalization Reagent stock solution by adding 100 μ L Protein Normalization Reconstitution Agent per tube. Thoroughly mix the reagent by pipetting 15 times.
- Use the following table to prepare the working solution of the reconstituted Protein Normalization Reagent stock solution. Thoroughly mix the working solution by pipetting 15 times.

PROTEIN MOLECULAR WEIGHT RANGE		
12–230 kDa		
LYSATE CONCENTRATION	Stock Solution	Reconstitution Agent
0.2–1.2 mg/mL	50 μ L	250 μ L

2. Detection Module preparation

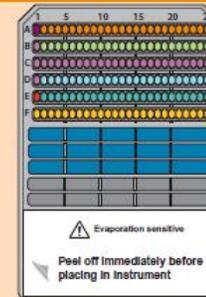
FOR CHEMILUMINESCENCE DETECTION

- Refer to Chemiluminescence Detection Module insert for instructions on primary and secondary antibody preparations

FOR FLUORESCENCE DETECTION

- Refer to Fluorescence Detection Module insert for instructions on primary and secondary antibody preparations

3. Pipette your plate

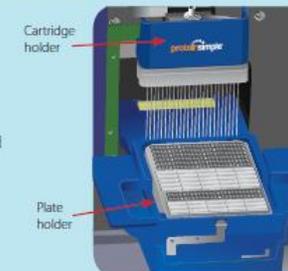


For more consistent results, keep the lid on the microplate between reagent additions and minimize bubble formation when adding Wash Buffer to the troughs and microplates. Protein Normalization reagent should be dispensed last.

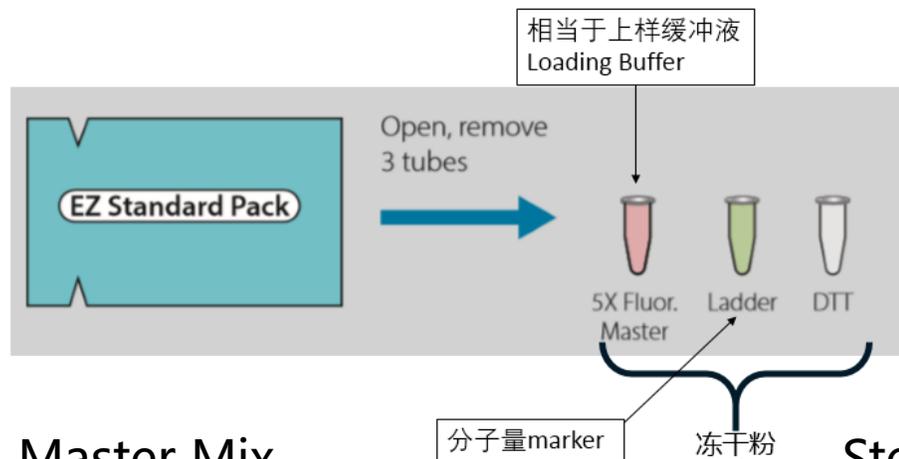
- Dispense reagents into the assay plate using the volumes shown in the plate diagram.
- Centrifuge the plate for 5 minutes at 2500 rpm (~1000 x g) at room temperature. Ensure liquid is fully down in all wells.

4. Start Jess

- Select the desired assay parameters in Compass software.
- Open Jess's door.
- Insert a capillary cartridge into the cartridge holder. The interior light will change from orange to blue.
- Remove the assay plate lid. Hold plate firmly on bench and carefully peel off evaporation seal. Pop any bubbles observed in the Separation Matrix wells with a pipette tip.
- Place the assay plate on the plate holder.
- Close Jess's door.
- Click the Start button in Compass.
- When the run is complete, discard the plate and cartridge.

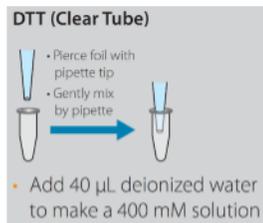
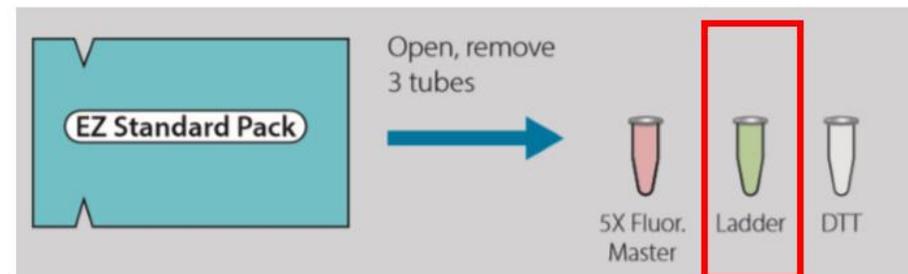
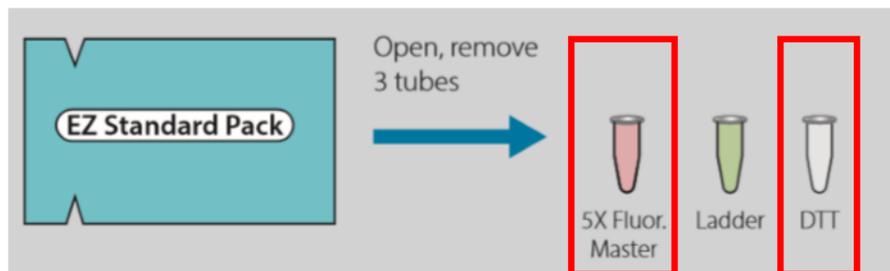


样品制备



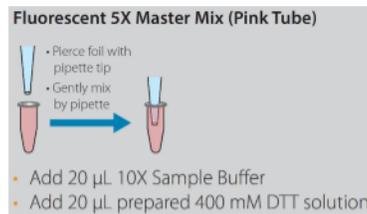
Step 1: 配制 5X FL Master Mix

Step 2: 配制分子量标准品 (marker)

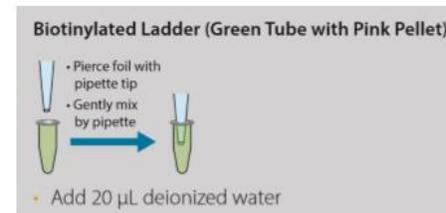


Reconstitute DTT

终浓度: 5x



Add: DTT
10x Sample Buffer



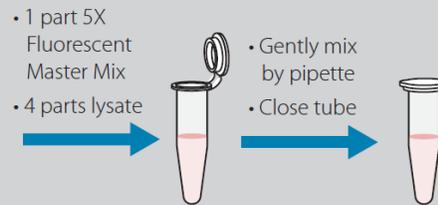
Add: 20 µL H₂O

样品制备

Step 3: 稀释样品

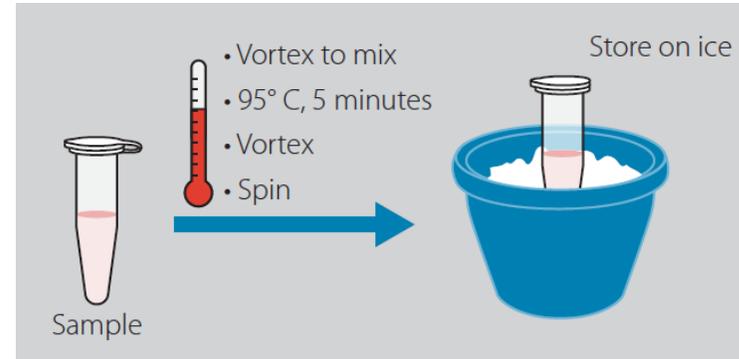
B PREPARE YOUR SAMPLES

- The optimal protein concentration depends on the expression level of your protein. Dilute lysate as necessary with 0.1X Sample Buffer.
- Combine 1 part 5X Fluorescent Master Mix with 4 parts diluted lysate in a microcentrifuge tube (final concentration 0.4 mg/mL for chemiluminescence or 1.0 mg/mL for fluorescence). Produce enough diluted sample volume required for assay.



样品用 **0.1X** Sample Buffer
(蛋白终浓度0.2-2mg/ml)

混合: **4** 体积 样品
1 体积 FL Master Mix



95°C 加热 5 分钟充分变性

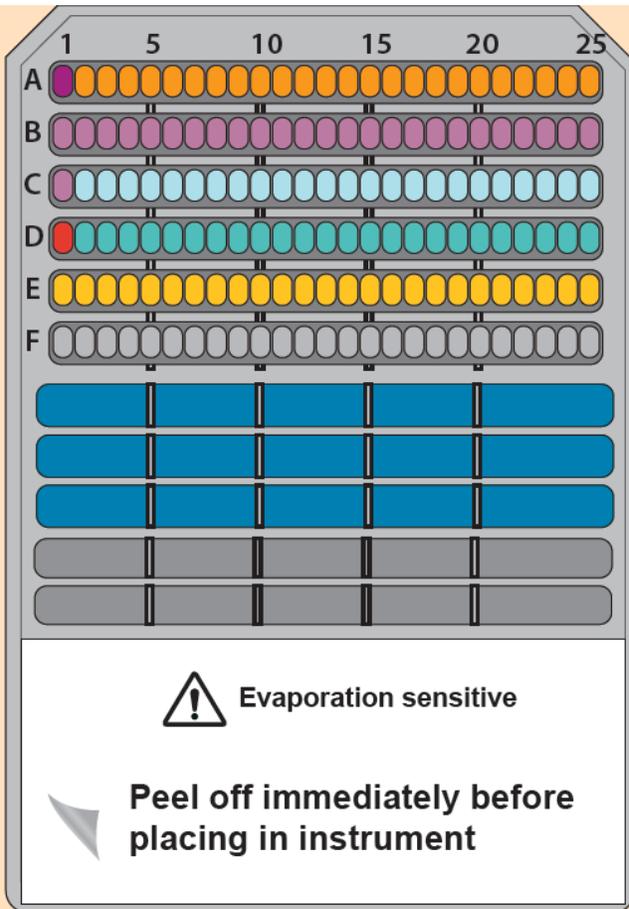
变性时建议用小管并封口以减少蒸发造成样品终浓度改变

- 需两次离心，以保证充分混合
 - ① 变性后先冰上冷却
 - ② Vortex混匀
 - ③ 短时离心
 - ④ Vortex混匀
 - ⑤ 短时离心

检查 Proteinsimple 提供的 “[Simple Western buffer compatibility table](#)”，确认裂解液兼容性

板布局 — 化学发光

第 1 个孔:
 分子量标准品 5 μL
 封闭液 10 μL
 封闭液 10 μL
 Streptavidin HRP
 发光底物 15 μL



■ Biotinylated Ladder, 5 μL ; ■ Prepared Samples, 3 μL
■ Wes Antibody Diluent 2, 10 μL
■ Wes Antibody Diluent 2, 10 μL ; ■ Primary Antibody, 10 μL
■ Streptavidin-HRP, 10 μL ; ■ Secondary HRP Conjugate, 10 μL
■ Luminol-Peroxide Mix, 15 μL

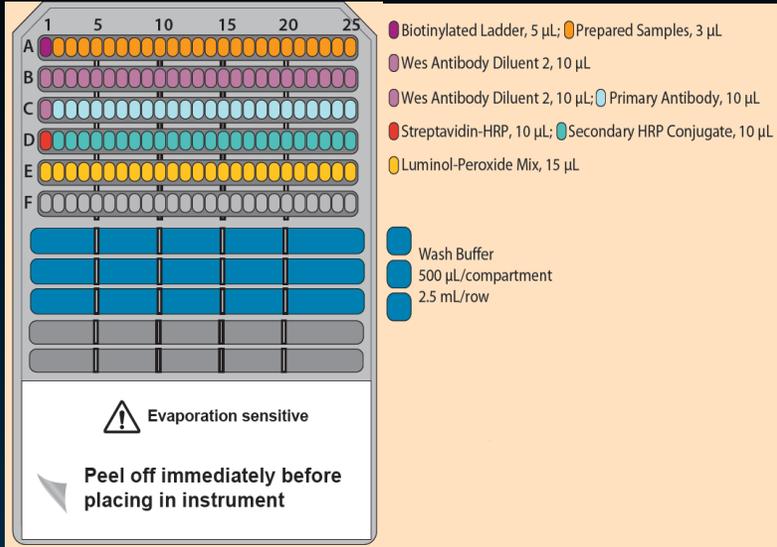
■ Wash Buffer
■ 500 μL /compartment
■ 2.5 mL/row

分离胶
 浓缩胶
 洗胶液
 电泳缓冲液
 电泳缓冲液

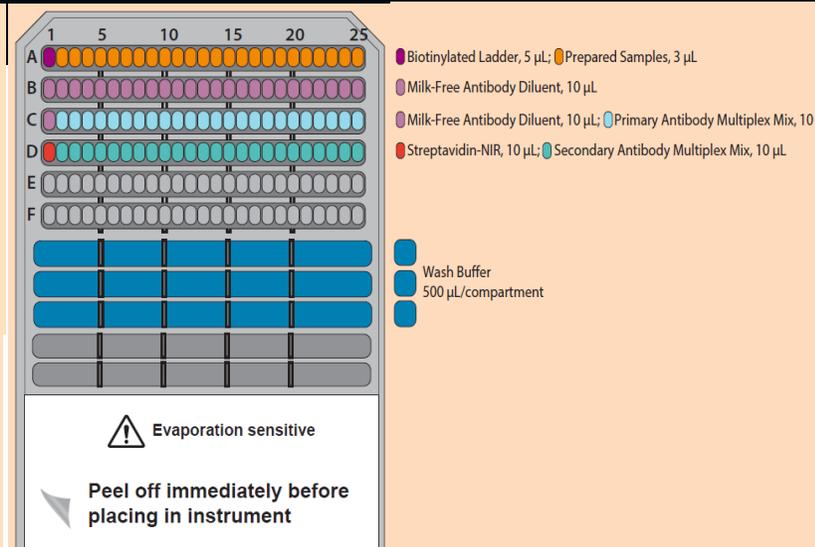
样本:
A: Sample 3 μL
B: 封闭液 10 μL
C: 稀释后的一抗 10 μL
D: 二抗 10 μL
E: 发光底物 15 μL

- 加样注意事项:
- ◆ 避免气泡: 反向吸液
 - ◆ 避免蒸发
 - 操作连贯
 - 加盖
 - 上机才撕膜
 - ◆ 离心: 室温, 加盖

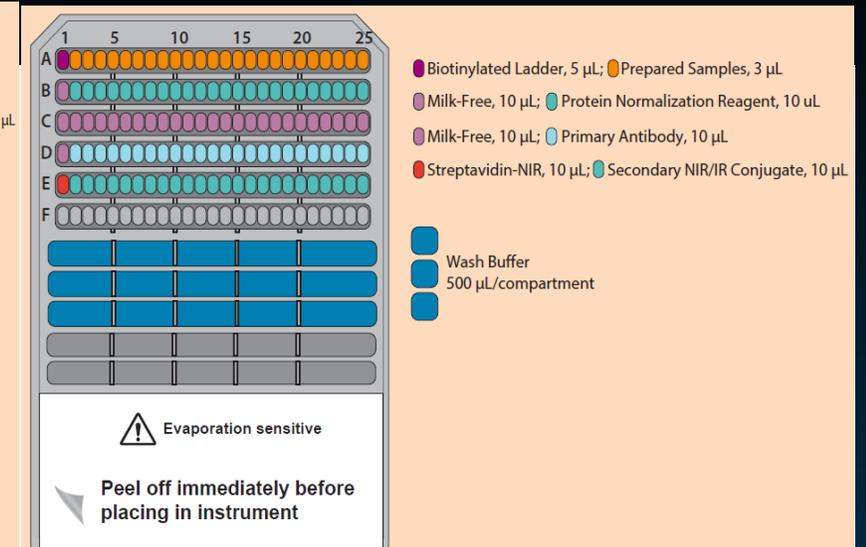
单独化学



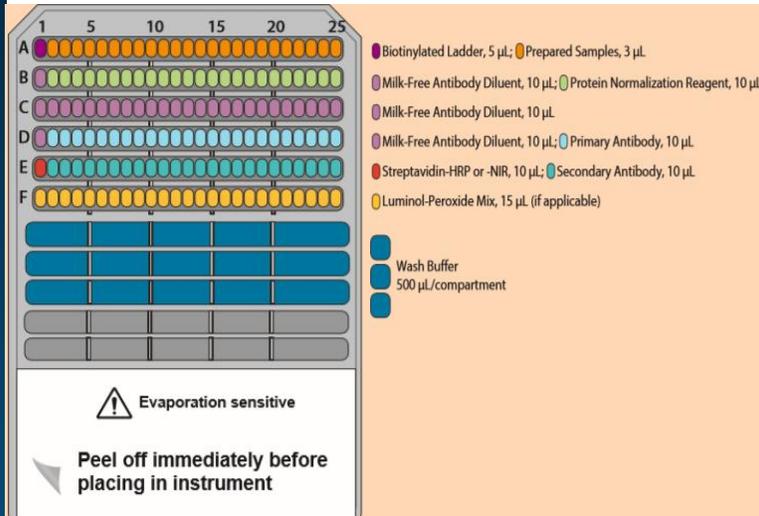
单独荧光



总蛋白和荧光



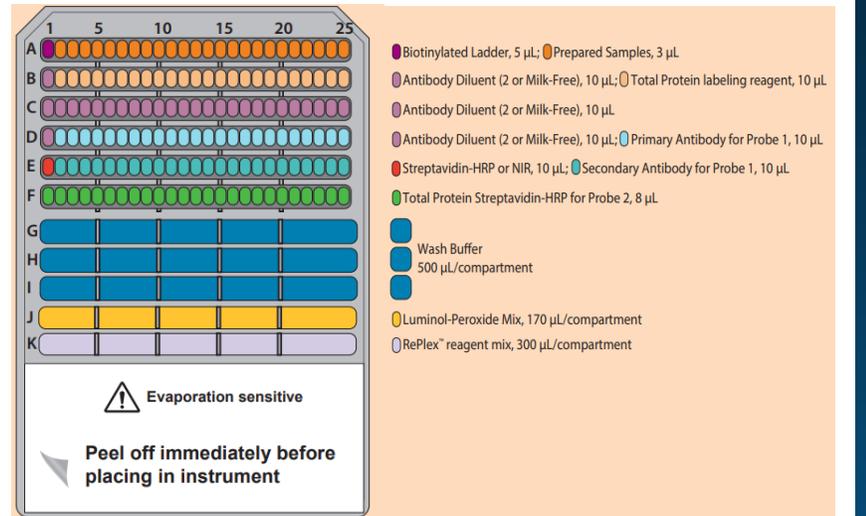
三者结合



两轮免疫



免疫和总蛋白



公众号报修流程



PROTEINSIMPLE SIMPLE YOUR PROTEIN ANALYSIS



4000-863-973



FluorChem
• Simple Imaging



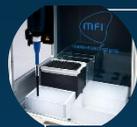
Jess/Wes/Abby
• Simple Western



Ella
• Simple ELISA



Milo
• Simple Sc-Western



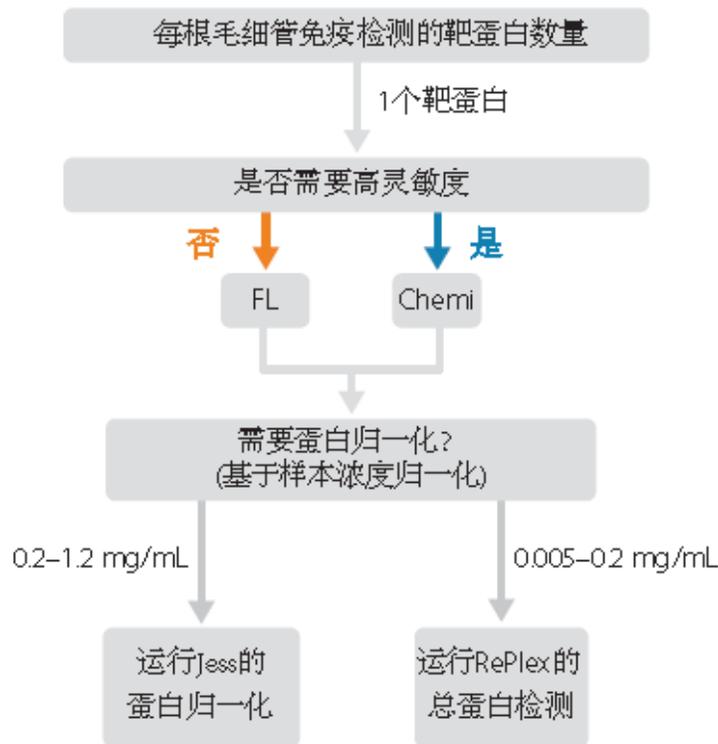
MFI
• Simple Particle Analysis



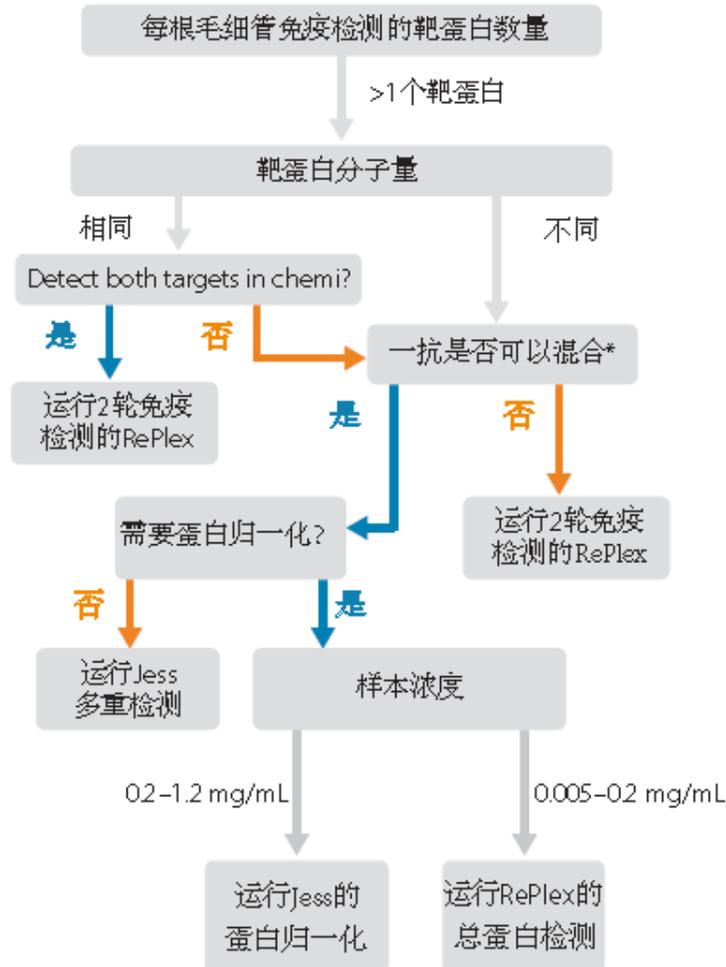
Maurice
• Simple icIEF + CE-SDS

JESS实验决策树

实验选择概述- 只检测1个靶蛋白



实验选择概述- 检测多于1个靶蛋白



RePlex: 如果第二轮进行化学发光检测, 推荐第一轮进行低丰度靶蛋白的化学发光或荧光检测。

多重检测: 多种一抗混在一起进行孵育后检测, 需验证可行性。

*检查将要混合在一起的抗体各自的交叉反应、动态范围以及背景。